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Title:
Perfect score:
Sequence:
                                                                                                                                                                                                                                                    Result
No.
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /SIDS8/gcgdat
2: /SIDS8/gcgdat
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2: /SIDSB/gcgdata/geneseq/geneseqp/AA1981.DAT:*
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6: /SIDSB/gcgdata/geneseq/geneseqp/AA1985.DAT:*
7: /SIDSB/gcgdata/geneseq/geneseqp/AA1987.DAT:*
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2000 Comp
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AAY28408
AAY92236
AAY92257
AAAY45062
AAB87601
AAB35260
AAB3526664
AAB35263
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AAB35263
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                                  Human IL-1 recepto
Human IL-1 homolog
Human TANGO-93 pro
Human PRO4342. Ho
Human IL-IL1. Hom
Interleukin-IL1 re
Protein encoded by
A human interleuki
                                                                                                                                                                                                                                            Description
                                                                                                                                                                                                       Human interleukin
Interleukin-IL1
Interleukin-IL1
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Sequence encoded b	AAP93616	10	165	38.8	319.5	5
4	AAB66670	22	159	•	319.5	14
_	AAB14829	21	159	•	319.5	Ω
acellular	AAW83008	20	159		319.5	12
IL-1 receptor anta	AAW73946	20	159	38.8	319.5	11
ro -	AAY28287	20	159	٠	319.5	0
	AAY33277	20	159		319.5	39
erved	AAY33275	20	159		319.5	8
Recombinant interl	AAW37787	19	159		319.5	37
Human intracellula	AAW80777	19	159		319.5	36
Variant Il-1 cytok	AAR15262	12	159	٠	319.5	35
\mathbf{r}	AAB51302	22	153		319.5	34
	AAW61149	19	153		319.5	3
þį	AAW22894	18	153		319.5	32
IL-1 inhibitor (IL	AAR35486	14	152	٠	319.5	3
Native IL-1ra poly	AAR27495	13	152		319.5	30
iCIL-1ra. Homo sa	AAR73642	16	159	39.4	324.5	29
	AAY97067	21	98	48.4	398	8
\vdash	AAY97068	21	94	51.0	420	27
Protein encoded by	AAB66663	22	80		437	26
A human interleuki	AAY43525	20	80	٠	437	25
en E	AAB35266	22	104		522.5	24
	AAY45061	21	156	89.2	734	23
መ	AAY92260	21	156		734	22
	AAY28407	20	156	•	734	21
	AAW86284	19	156		734	20
Murine interleukin	AAB48828	22	155		734	9
ion relate	AAB66672	22	155		734	8
II-II	AAB35261	22	155		734	17
IL-1 recep	AAY96937	21	155		734	6
human IL-	AAY92255	21	155		805	5
c huma	AAY92253	21	155		805	14
n IL-1	AAY92254	21	155	98.7	812	<u>_</u> 3
Human IL-1 homolog	AAY92256	21	155	99.1	816	12

ALIGNMENTS

RESULT AAY28408

AAY28408 standard; Protein;

155

B

Human interleukin 1 delta

28-SEP-1999 AAY28408;

(first entry)

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PPTX RRAPEX RPH X RRAPEX RRAPE
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Murine and Human interleukin 1 delta DNA, polypeptides and its fragments, useful as molecular weight markers
                                                                                                                                                       WPI; 1999-458310/38.
N-PSDB; AAX89432.
                                                                                                                                                                                                                                                                                                          Sims JE;
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09-JAN-1998;
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98US-0071074
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Best Local
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                                                                              23-DEC-1998;
22-JAN-1999;
13-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               isoelectric point of a sample protein. Antibodies generated against IL-1 delta and its fragmented peptides can be used to enhance the accuracy of these molecular weight markers to determine the apparent molecular eight and isoelectric point of a sample protein. IL-1 delta can be used to screen for potential inhibitors of activity associated with IL-1 delta counter structure molecules. IL-1 delta can also be used as therapeutic agents for the treatment of diseases mediated by IL-1 delta. II-1 delta may be used as a reagent in studying the interleukin 1 (IL-1) signalling pathway, or as a reagent to block IL-1 signalling. The IL-1 delta coding sequences can be used to identify human chromosome 2, and to identify genes associated with certain diseases, especially with region 2q11-12, including glaucoma, ectodermal dysplasia, insulindependent diabetes mellitus, wrinkly skin syndrome, T-cell leukemia/
             Goddard A,
                                                                                                                                                22-DEC-1999;
                                                                                                                                                                                                               WO200039297-A2
                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                             hIL-TRa3; human interleukin-1 receptor antagonist-3; IL-11p; osteopathic; interleukin-1-like polypeptide; anti-inflammatory; amti-asthmatic; anti-arthritic; antinicrobial; respiratory; anti-ischemic; vaccine; dermatalogical; immunomodulatory; gastrointestinal; gene therapy.
                                                                                                                                                                                                                                                                                                                                                             Human IL-1 receptor antagonist
                                                                                                                                                                                                                                                                                                                                                                                              31-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY96936 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       weight of a sample protein. The protein and its fragments are useful as controls for peptide fragmentation. This is useful for determining the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1;
                                            (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          lymphoma and tibial muscular dystrophy.
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1 delta proteins are useful for the determination of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LCTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD 155
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155; Conserv
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                                                                             98US-0113430.
99US-0116843.
99US-0129122.
                                                                                                                                                99WO-US30720
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Protein;
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Pred. No. 1.2e-86;
); Mismatches 0;
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RESULT
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                                                                                                                                                                                                   Generic; interleukin-1; IL-1; homologue; zilla3; anti-inflammatory; antiagonist; pro-inflammatory; agonist; immunomodulator; antiarthritic; antirheumatic; osteopathic; antipsoriatic; antibacterial; cytostatic; immunosuppressive; antiulcer; antidiabetic; nephrotropic; vasotropic;
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                                                                                                                                                                                                                                                                                                                                                                                            AAY92257 standard; Protein; 155
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N-PSDB; AAA51597.
Peptide
                               Peptide
                                                                 Peptide
                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                          Human IL-1 homologue, hzilla3.
                                                                                                                                                                                                                                                                                                                            10-AUG-2000
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                                 /label= loop
17. 21
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14..16
                                                                                                                   Location/Qualifiers
              'label= beta-strand
                                                                                 beta-strand
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Pred. No. 1.2e-86;
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This shows an interleukin-1 (IL-1) homologue, designated zilla: It is believed that zilla3 acts through IL-1 receptors. In ger zilla3 proteins having a Lys residue at position 148 will have anti-inflammatory activity (e.g. AAY92256), whilst those having
                                                   Proteins useful for treatment of inflammatory conditions such as rheumatoid arthritis and psoriasis are agonists or antagonists forms new interleukin-1 homologue
                                        Disclosure;
                                                                                                                    (ZYMO ) ZYMOGENETICS INC
                                                                                                                                    08-OCT-1998;
                                                                                                                                                  08-OCT-1999;
                                                                                                                                                                               WO200020595-A1
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DB; AAA09193, AAA09194.
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                                       Page
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26..28
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146..14
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72..76
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117..121
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Best Local S
Matches 155
                                                                                                                                                                                                                                                                                                                                                                                          TANGO-93; cytokine; human; secreted protein; II-1 expression; cancer; Interleukin-1 receptor antagonist; II-1ra; inflammation; antiasthmatic; immunosuppressive; antirheumatic; antiarthritic; antipsoriatic; asthma; antiinflammatory; antibacterial; antiulcer; cytostatic; immunomodulator; osteopathic; dermatological; antidiabetic; psoriasis; ulcerative colitis; graft vs-host disease; rheumatoid arthritis; inflammatory bowel disease; septic shock; cachexia; Crohn's disease; chronic myelogenous leukaemia; liver disease; diabetes; osteopathit; Hodgkin's disease; Lyme disease; liver disease; diabetes; osteopathitis; Hodgkin's disease; Lyme disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (see AAY92254) or Glu at this position will have pro-inflammatory action. Zilla3 is used to modulate an immune response in an animal (claimed). Antagonists zilla3 forms may be used to treat or prevent chronic inflammatory diseases such as rheumatoid arthritis, osteoarthritis and Lyme arthritis, psoriasis, to reduce tissue damage after ischemia, to treat septic shock, graft-versus-host disease and leukemia. The antagonists may also alleviate inflammatory bowel disease including Crohn's disease and ulcerative collits, insulin-dependent including Crohn's disease and ulcerative collits.
                                                                                                                                                                                                                                                                                                                                                           autoimmune disease; myasthenia gravis; pharmacogenomic;
chromosome 2; systemic lupus erythematosus; forensic; t;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diabetes mellitus, acute pancreatitis, glomerulonephritis and ischemia. Agonist forms of zilla3 may promote wound healing by effects on growth factor secretion and cell proliferation. The treat infections, especially gastrointestinal infections.
                                                                                                                                                                                                                 06-AUG-1999;
                                                                                                                                                                               07-AUG-1998;
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No. 1.2e-86;
matches 0;
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WPI; 200 N-PSDB;

2000-205669/18. DB; AAZ50812.

Isolated nucleic acid sequences encoding TANGO-93 polypeptide useful for treating a variety of cellular processes e.g. asthma, rheumatoid

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ID ACCOMPAND ACC
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Best Local
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09-DEC-1999;
11-JAN-2000;
18-FEB-2000;
18-FEB-2000;
22-FEB-2000;
01-MAR-2000;
03-MAR-2000;
25-APR-2000;
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15-SEP-1999;
07-DEC-1999;
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2000US-0175481.
2000WO-US04341.
2000WO-US04342.
2000WO-US04414.
2000WO-US05601.
2000US-0187202.
2000US-019397.
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99WO-US21090.
99US-0169495.
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Pred. No. 1.2e-86;
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Best Local Similarity
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Grimaldi CJ,
                                                                                                                                                                                                                                                                                                                                                                     Human; IL-II1; interleukin-1 locus; IL-lbeta; IL-lreceptor; chromosome 2q13; inflammatory disease; heart disease; Graves rheumatoid arthritis; inflammatory bowel disorder; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 12; Fig 152; 278pp; English.
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05-JUN-2000; 2000US-0209832.
                                                                           16-JUL-1999;
                                                                                                                                                                                                                                       W0200105974-A2
                                                                                                                                                                                                                                                                                                                                            osteoporosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human IL-IL1.
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                                                                                                                             17-JUL-2000; 2000WO-US19508
                                                                                                                                                                                                                                                                                            Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LCTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              lctvpeadqpvrltqlpenggwnapitdfyfqqcd
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            155 AA;
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Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                 systemic lupus erythematosus
                                                                           99US-0144298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protein;
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Watanabe
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Pred. No. 1.2e-86;
); Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRO polypeptides, useful ir s hybridization probes, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ç (
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sis. The PRO
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(INTE-) INTERLEUKIN GENETICS INC

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RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 155; Conser
         Nucleic acids encoding human and murine interleukin-111 polypeptides useful for controlling inflammatory processes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention provides the protein and coding sequences of the human and murine interleukin-IL1 (IL-IL1) proteins. The IL-IL1 gene is located between the IL-lbeta and IL-lreceptor genes at human chromosome 2q13. The sequences are useful in the diagnosis, prevention and treatment of heart disease, cancer and inflammatory diseases such as rheumatoid arthritis, systemic lupus erythematosus, inflammatory bowel disorder, diabetes, psoriasis, osteoporosis, lichen sclerosis, ulcerative colitis, severe periodontal disease and pregnancy complications. The present sequence is the human IL-IL1 protein.
                                                                                                                                                                                                              WO200105974-A2
                                                                                                                                                                                                                                      Unidentified
                                                                                                                                                                                                                                                             Mouse; IL-ILI; interleukin-1 locus; IL-lbeta; IL-lreceptor; psoriasis; chromosome 2q13; inflammatory disease; heart disease; Graves' disease; rheumatoid arthritis; inflammatory bowel disorder; diabetes; cancer; osteoporosis; systemic lupus erythematosus; human.
                                                                                                                                16-JUL-1999;
                                                                                                                                                      17-JUL-2000; 2000WO-US19508
                                                                                                                                                                                                                                                                                                                                                            08-MAY-2001
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                                                                                                      (INTE-) INTERLEUKIN GENETICS
                                                                                                                                                                                     25-JAN-2001
                                                                                                                                                                                                                                                                                                                               Interleukin-IL1 recombinant protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 vqggsqclscgvgqeptltlepvnimelylgakesksftfyrrdmgltssfesaaypgwf
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VQGGSQCLSCGVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            controlling inflammatory processes
                                                                             Barton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; ilarity 100.0%; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                               Protein;
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Pred. No. 1.2e-86;
; Mismatches 0;
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Best Local
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                                                                                                                                                                                                   13-OCT-1999;
08-DEC-1999;
10-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention provides the protein and coding sequences of the human and murine interleukin-III (II-III) proteins. The II-III gene is located between the II-lbeta and II-lreceptor genes at human chromosome 2q13. The sequences are useful in the diagnosis, prevention and treatment of heart disease, cancer and inflammatory diseases such as rheumatoid arthritis, systemic lupus erythematosus, inflammatory bowel disorder, diabetes, psoriasis, osteoporosis, lichen sclerosis, ulcerative colitis, severe periodontal disease and pregnancy complications. The present
                                 Claim 1;
                                                       Isolated nucleic acids encoding interleukin-1 (IL-1) antagonist proteins (referred as IL-1Hy1), useful in cancer, e.g. breast adenocarcinoma and brain tumors, inflammatory disease mediated by IL-18
                                                                                                                                           Ford
                                                                                                                                                                                                                                       07-JUL-1999;
                                                                                                                                                                                                                                                              07-JUL-2000; 2000WO-US18710
                                                                                                                                                                                                                                                                                                               WO200102571-A2
                                                                                                                                                                                                                                                                                                                                                                                    Protein encoded by extended
                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB66664;
                                                                                                                   WPI; 2001-071582/08.
                                                                                                                                                                  (HYSE-) HYSEQ
                                                                                                                                                                                                                                                                                                                                                            interleukin; IL-1 receptor; cancer; inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB66664 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence is a recombinant IL-IL1 protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          VQGGSQCLSCGVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                Fig 6; 179pp; English.
                                                                                                                                           Pace
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fig 6; 150pp;
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99US-0457626.
2000US-0523552.
2000US-0576008.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                                                  INC.
                                                                                                                                                                                                                                         99US-0348942
                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein;
                                                                                                                                                                                                                                                                                                                                                                                                           entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        English.
                                                                                                                                                                                                                                                                                                                                                                                  B2HFLS20W cDNA library sequence #2
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                                                                  the treatment and an
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The present invention relates to interleukin (IL)-1 antagonist proteins. IL-1Hyl is useful for treating

cancer receptor

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RESULT
AAY43526
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Best Local
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                                                                                                                                                                03-APR-1998;
15-MAY-1998;
20-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           an inflammatory disease mediated by IL-18, inflammation resulting from infection or allergic reactions, and inflammation associated with chronic bronchitis, arthritis, diabetes or
New isolated interleukin-1 receptor binding polypeptides, used
                                                          Drmanac R,
Leshkowitz
                                                                                                                               31-JUL-1998;
13-JAN-1999;
                                                                                                                                                                                                               05-APR-1999;
                                                                                                                                                                                                                                      14-OCT-1999.
                                                                                                                                                                                                                                                            W09951744-A2
                                                                                                                                                                                                                                                                                                        anti-inflammatory
                                                                                                                                                                                                                                                                                                                  hematopoiesis regulating activity; tissue growth activity; activin activity; inhibin activity; chemotactic activity; chemokinetic activity; hemostatic activity; thrombolytic activity;
                                                                                                                                                                                                                                                                                                                                                    graft versus host disease; inflammatory bowel disease; inflammation; pulmonary disease; autoimmune disease; inflammatory disease; antiproliferative; myelogenous leukemia; premature labor; intrauterine infection; nutritional activity;
                                                                                                                                                                                                                                                                                                                                                                                                   Human; interleukin-1 receptor; IL-1; antagonist; sepsis; acute pancreatitis; endotoxic shock; cytokine induced shock; rheumatoid arthritis; chronic inflammatory arthritis; pancreatic cell damage; diabetes mellitus type 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              A human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAY43526 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    endothermia.
                                                                                           (HYSE-)
                                                                                                                                                      19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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                         AAZ30050
                                                                                                                                                                                                                                                                                                                                                                                                                                                            interleukin-1 receptor antagonist.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              155 AA;
                                                        Crkvenjakov
), Kita D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                             INC.
                                                                                                                  98US-0099818.
98US-0127698.
99US-0229591.
99US-0251370.
                                                                                                                                                                98US-0055010.
98US-0079909.
98US-0082364.
                                                                                                                                                                                                               99WO-US04291
                                                                                                                                                                                                                                                                                                        activity.
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                                                          Ford
                                                         Dickson M,
d J, Pace /
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 823; DB 22;
Pred. No. 1.2e-86;
; Mismatches 0;
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                                                         Drmanac S,
, Alfenito
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 155;
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                                                                     Labat
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to treat
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AAB35263
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches 154;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         arthritis, chronic inflammatory arthritis, pancreatic cell damage from diabetes mellitus type 1, graft versus host disease, inflammatory bowel disease, inflammatory activity, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents a human interleukin-1 (IL-1) receptor antagonist. The polypeptide is capable of binding IL-1 receptors (IL-1Rs). The polypucleotides and polypeptides can be us the prevention or treatment of disorders involving sepsis, acute pancreatitis, endotoxic shock, cytokine induced shock, rheumatoid
                      Nicklin M,
                                                                                                                                                                                                                                                                                                                                                                                                              osteoporosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                     chromosome 2q13; inflammatory disease; heart disease; Graves' diseas rheumatoid arthritis; inflammatory bowel disorder; diabetes; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hemostatic and thrombolytic activity, receptor/ligand activity, and anti-inflammatory activity. The products can also be used for detection, diagnosis and drug screening.
                                                                          (INTE-) INTERLEUKIN GENETICS INC
                                                                                                                                                                                       17-JUL-2000; 2000WO-US19508
                                                                                                                                                                                                                                                                                                    WO200105974-A2
                                                                                                                                                                                                                                                                                                                                                          Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse; IL-IL1; interleukin-1 locus; IL-lbeta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Interleukin-IL1 recombinant protein #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB35263 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             e.g. sepsis, shock, arthritis, pancreatitis, graft-versus-host disease, inflammatory disease, autoimmune disease or proliferative disease
                                                                                                                                16-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LCTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VQGGSQCLSCGVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mvlsgalcfrmkdsalkvlylhnnqllagglhagkvikgeeisvvpnrwldaslspvilg 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                155 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6; 123pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                           systemic lupus erythematosus; human.
                                                                                                                                99US-0144298
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 820; DB 20; Pred. No. 2.7e-86;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IL-lreceptor; psoriasis;
lisease; Graves' disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  can be used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 154;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse; IL-IL1; interleukin-1 locus; IL-1beta; IL-1receptor; psoriasi chromosome 2q13; inflammatory disease; heart disease; Graves' disease rheumatoid arthritis; inflammatory bowel disorder; diabetes; cancer; osteoporosis; systemic lupus erythematosus; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of heart disease, cancer and inflammatory diseases such as rheumatoid arthritis, systemic lupus erythematosus, inflammatory bowel disorder, diabetes, psoriasis, osteoporosis, lichen sclerosis, ulcerative colitis, severe periodontal disease and pregnancy complications. The present sequence is a recombinant II-ILl protein.
                                                                                                                                                                                                                       Nicklin M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention provides the protein and coding sequences of the human and murine interleukin-IL1 (IL-IL1) proteins. The IL-IL1 gene is located between the IL-lbeta and IL-laceptor genes at human chromosome 2013. The sequences are useful in the diagnosis, prevention and treatment
                          Examples;
                                                                            Nucleic acids encoding useful for controlling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB35264;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAB35264 standard; Protein;
                                                                                                                                                                                                                                                                             (INTE-) INTERLEUKIN GENETICS
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                                                                                                                                                                                                                                                                                                                                                                                             17-JUL-2000; 2000WO-US19508
                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200105974-A2
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                                                                                                                                                                     2001-091974/10.
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for controlling inflammatory processes -
                          Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fig 6; 150pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            154 AA;
                                                                                                                                                                                                                          Barton J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                          6,
                                                                                                                                                                                                                                                                                                                                        99US-0144298.
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                                                                         human and murine interleukin-1L1 polypeptides inflammatory processes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IL-lreceptor; psoriasis;
lisease; Graves' disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention provides the protein and coding sequences of the human and murine interleukin-IL1 (IL-IL1) proteins. The IL-IL1 gene is located between the IL-lbeta and IL-Irceptor genes at human chromosome 2q13. The sequences are useful in the diagnosis, prevention and treatmen of heart disease, cancer and inflammatory diseases such as rheumatoid arthritis, systemic lupus erythematosus, inflammatory bowel disorder, diabetes, psoriasis, osteoporosis, lichen sclerosis, ulcerative colitis, severe periodontal disease and pregnancy complications. The present sequence is a recombinant IL-IL1 protein.
This polypeptide is variant interleukin-1 (IL-1) homologue, designated zilla3-K148. It is believed that zilla3 acts through IL-1 receptors. In general, zilla3 proteins having a Lys residue at position 148 will have anti-inflammatory activity, whilst those having Asp (see AAY92254)
                                                                                                                                                                                                                                                                                                                                                                                                           antagonist; pro-inflammatory; agonist; immunomodulator; antiarthritic;
antirheumatic; osteopathic; antipsoriatic; antibacterial; cytostatic;
immunosuppressive; antiulcer; antidiabetic; nephrotropic; vasotropic;
                                                                       Claim
                                                                                                               Proteins useful for treatment of inflammatory conditions such as rheumatoid arthritis and psoriasis are agonists or antagonists forms
                                                                                                                                                                                                Sheppard PO,
                                                                                                                                                                                                                                                        08-OCT-1998;
                                                                                                                                                                                                                                                                                  08-OCT-1999;
                                                                                                                                                                                                                                                                                                              13-APR-2000
                                                                                                                                                                                                                                                                                                                                          WO200020595-A1
                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                vulnerary.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Generic; interleukin-1; IL-1; homologue; zilla3; anti-inflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY92256;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY92256 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                 new interleukin-1 homologue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-AUG-2000 (first entry)
                                                                                                                                                                                                                           (ZYMO ) ZYMOGENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ctvpeadqpvrltqlpenggwnapitdfyfqqcd
                                                                                                                                                                      2000-303780/26
                                                                       4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IL-1 homologue, hzilla3-K148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             154;
                                                                                                                                                         AAA09195
                                                                       Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                      56-57; 64pp;
                                                                                                                                                                                                West RR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AA;
                                                                                                                                                                                                                                                        98US-0169745
                                                                                                                                                                                                                                                                                    99WO-US23533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein;
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100.0%;
                                                                                                                                                                                                Clegg
                                                                      English.
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RESULT
AAY92254
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or Glu at this position will have pro-inflammatory action. 2illa3 is used to modulate an immune response in an animal (claimed). Antagonists zilla3 forms may be used to treat or prevent chronic inflammatory diseases such as rheumatoid arthritis, osteoarthritis and lyme arthritis, psoriasis, to reduce tissue damage after ischemia, to treat septic shock, graft-versus-host disease and leukemia. The antagonists may also alleviate inflammatory bowel disease including Crohn's disease and ulcerative colitis, insulin-dependent diabetes mellitus, acute pancreatitis, glomerulonephritis and cerebral ischemia. Agonist forms of zilla3 may promote wound healing by II-1 effects on growth factor secretion and cell proliferation. They may also treat
Proteins useful for treatment of inflammatory conditions such as rheumatoid arthritis and psoriasis are agonists or antagonists fr
                                                                      Sheppard PO,
                                                                                                                                                        08-OCT-1999;
                                                                                                                                                                                                                                                         Misc-difference 99
                                                                                                                                                                                                                                                                                                                                                                  antagonist; pro'inflammatory; agonist; immunomodulator; antiarthritic;
antirheumatic; osteopathic; antipsoriatic; antibacterial; cytostatic;
immunosuppressive; antiulcer; antidiabetic; nephrotropic; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                             Generic;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Hhuman IL-1 homologue, hzilla3-D148.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY92254 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                    13-APR-2000
                                                                                                                                                                                                                WO200020595-A1
                                                                                                                                                                                                                                                                                   Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-AUG-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61
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                                                                                                                                                                                                                                                                                                                           sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LCTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                vqggsqclscgvgqeptltlepvnimelylgakesksftfyrrdmgltssfesaaypgwf 120
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                                                                                                                                                                                                                                                                                                                                                                                                             interleukin-1; IL-1; homologue; zilla3; anti-inflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         155 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                    West RR,
                                                                                                                            98US-0169745
                                                                                                                                                        99WO-US23533
                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                          /label= Ala, Ile, Thr
                                                                                                                                                                                                                                                                   /label= Glu, Lys
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein;
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Pred. No. 7.6e-86;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chronic inflammatory diseases such as rheumatoid arthritis, costeoarthritis and Lyme arthritis, psoriasis, to reduce tissue damage after ischemia, to treat septic shock, graft-versus-host disease and leukemia. The antagonists may also alleviate inflammatory bowel disease including Crohn's disease and ulcerative collitis, insulin-dependent diabetes mellitus, acute pancreatitis, glomerulonephritis and cerebral ischemia. Agonist forms of zilla3 may promote wound healing by IL-1 effects on growth factor secretion and cell proliferation. They may also treat infections, especially gastrointestinal infections. Note: This sequence is not given in the specification, it is created from SEQ. ID. 7, which is given on pages 55-56.
                                                                                                                                                                                                                                                Generic; interleukin-1; IL-1; homologue; zilla3; anti-inflammatory; antagonist; pro-inflammatory; agonist; immunomodulator; antiarthritic; antirheumatic; osteopathic; antipsoriatic; antibacterial; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This is a generic interleukin-1 (IL-1) homologue, designated zilla3. It is believed that zilla3 acts through IL-1 receptors. In general, zilla3 proteins having a Lys residue at position 148 (see AAY92255 and AAY92256) will have anti-inflammatory activity, whilst those having Asp (i.e. this sequence) or Glu at this position will have pro-inflammatory action. Zilla3 is used to modulate an immune response in an animal (claimed). Antagonists zilla3 forms may be used to treat or prevent
                 WO200020595-A1
                                                                    Misc-difference
                                                                                                    Misc-difference
                                                                                                                                                                                                                                                                                                                        Generic human
                                                                                                                                                                                                                                                                                                                                                       10-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                          AAY92253 standard; protein; 155 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                 Misc-difference
                                                                                                                                                                                    Homo sapiens.
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                                                                                                                                                                                                                                    mmunosuppressive; antiulcer; antidiabetic; nephrotropic; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           lctvpeadqpvrltqlpenggwnapitdfyfqqcd 155
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mvlsgalcfrmkdsalkvlylhnnqllagglhagkvikgexisvvpnrwldaslspvilg 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                        IL-1
                                                                /label= Ala,
148
                                                                                                                                                   Location/Qualifiers
                                                 /label=
                                                                                                                  'label=
                                                                                                                                                                                                                                                                                                                      homologue, hzilla3-X148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98.7%;
98.7%;
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                                                 Glu,
                                                                                                                  Glu,
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                                                 Lys, Asp
                                                                                  Ile,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 812; DB 21;
Pred. No. 2.2e-85;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 155;
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                                                                                                                                                                                                                                                                                                                                              RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local !
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 152;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             zilla3. It is believed that zilla3 acts through IL-1 receptors. In general, zilla3 proteins having a Lys residue at position 148 (see AAY92255 will have anti-inflammatory activity, whilst those having Asp (see AAY92254) or Glu at this position will have pro-inflammatory action. Zilla3 is used to modulate an immune response in an animal (claimed). Antagonists zilla3 forms may be used to treat or prevent chronic inflammatory diseases such as rheumatoid arthritis, osteoarthritis and Lyme arthritis, psoriasis, to reduce tissue damage after ischamia, to treat septic shock, graft-versus-host disease and leukemia. The antagonists may also alleviate inflammatory bowel disease including Crohn's disease and ulcerative collits, insulin-dependent disease and linear actions and core beautiful to the contraction of the collision of the collisions and core beautiful disease and college of the collisions and core beautiful disease and college of the collisions.
 Misc-difference
                                                                                                     Generic; interleukin-1; IL-1; homologue; zilla3; anti-inflammatory; antagonist; pro-inflammatory; agonist; immunomodulator; antiarthritic; antiheumatic; osteopathic; antipsoriatic; antibacteria1; cytostatic; immunosuppressive; antiulcer; antidiabetic; nephrotropic; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diabetes mellitus, acute pancreatitis, glomerulonephritis and cerebral ischemia. Agonist forms of zilla3 may promote wound healing by IL-1 effects on growth factor secretion and cell proliferation. They may a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Proteins useful for treatment of inflammatory conditions such as rheumatoid arthritis and psoriasis are agonists or antagonists forms of
                                                                                                                                                                                                 Generic human IL-1 homologue,
                                                                                                                                                                                                                                       10-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 55-56; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sheppard PO,
                                                      Homo sapiens
                                                                                          vulnerary.
                                                                                                                                                                                                                                                                                                          AAY92255 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treat infections, especially gastrointestinal infections.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ZYMO ) ZYMOGENETICS INC
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                                                                                                                                                                                                                                                                                                                                                                                                                 121 LCTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD 155
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                                                                                                                                                                                                                                                                                                                                                                                               lctvpeadqpvrltqlpenggwnapitxfyfqqcd 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        vqggsqclscgvgqeptltlepvnimelylgakesksfxfyrrdmgltssfesaaypgwf 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    polypeptide is a generic interleukin-1 (IL-1) homologue, designated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98US-0169745
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   Location/Qualifiers
                                                                                                                                                                                                                                                                                                          protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97.8%;
98.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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                                                                                                                                                                                                   hzilla3-K148
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 155;
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Search completed: Job time: 27 sec

June

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2001, 14:05:25

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CC This is a generic interleukin-1 (II-1) homologue, designated zilla3. CC It is believed that zilla3 acts through II-1 receptors. In general, CC zilla3 proteins having a Lys residue at position 188 (this sequence CC and AAY92256) will have anti-inflammatory activity, whilst those having Asp (AAY92254) or Glu at this position will have pro-inflammatory CC action. Zilla3 is used to modulate an immune response in an animal CC (claimed). Antagonists zilla3 forms may be used to treat or prevent CC chronic inflammatory diseases such as rheumatoid arthritis, contect including Crohn's diseases such as rheumatoid arthritis, of the antagonists may also alleviate inflammatory bowel disease and CC leukemia. The antagonists may also alleviate inflammatory bowel disease including Crohn's disease and ulcerative colitis, insulin-dependent CC diabetes mellitus, acute pancreatitis, glomerulonephritis and cerebral conscience is a growth factor secretion and cell proliferation. They may CC also treat infections, especially gastrointestinal infections.

CC Note: This sequence is not given in the specification, it is created. CC from SEQ. ID. 7, which is given on pages 55-56.
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                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 3; Page -; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sheppard PO,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  new interleukin-1 homologue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ZYMO ) ZYMOGENETICS INC
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    121
                                121
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                                                                                                                                                                                                        Local Similarity
                                                                                                                      vqggsqclscgvgqeptltlepvnimelylgakesksfxfyrrdmgltssfesaaypgwf
                                                                                           VQGGSQCLSCGVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 120
                                                                                                                                                                                                                                                                       155 AA;
                                                                                                                                                                                         Conservative
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Pred. No. 1.4e-84;
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methionyl-tRNA syn
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9	69.5	70	70	70	70.5	70.5	71	71	71	71.5	71.5	71.5	71.5	71.5	. 72
6.4	8.4	8.5	8.5	8 . 5	8.6	8.6	8.6	8.6	8.6	8.7	8.7	8.7	8.7	8.7	8.7
152	801	323	305	269	1161	778	611	535	475	1110	1036	700	578	551	423
κ.	N	N	N	N	N	N	N	N	N	N	N	N	N	N	N
B43863	B86673	A82433	E83091	C83516	T18642	D72421	A48582	T06285	G84697	T33877	B83466	T34321	T45107	H72117	в86214
hypothetical prote	penicillin-binding	fructokinase VCA06	ATP sulfurylase sm		hypothetical prote	xylosidase - Therm	vacuolar ATPase A	_	hypothetical prote	hypothetical prote	•	hypothetical prote	H+-transporting AT	methioninetRNA l	hypothetical prote

ALIGNMENTS

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interleukin-1 receptor antagonist secreted form precursor - human (;Species: Homo sapiens (man) (;Species: Homo sapiens (man) (;Species: O7-Jun-1990 #sequence_revision 07-Jun-1990 #text_change 26-May-2000 (;Accession: A40956; 137894; A30368; S08160; S08159; A37822 R;Eisenberg, S.P.; Brewer, M.T.; Verderber, E.; Heimdal, P.; Brandhuber, B.J.; The Proc. Natl. Acad. Sci. U.S.A. 88, 5232-5236, 1991 A;Title: Interleukin 1 receptor antagonist is a member of the interleukin 1 gene: A,Reference number: A40956; MUID:91271363 A;Accession: A40956; MUID:91271363 A;Accession: A40956 A;Gross-references: GB:M63099; NID:9186385; PIDN:AAB41943.1; PID:9186386 A;Cross-references: GB:M63099; NID:9186385; PIDN:AAB41943.1; PID:9186385 A;Cross-r
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A;Map position: 2q14
C;Keywords: macrophage
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C;Species: Homo sapiens (man)
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 21-Ju1-2000
C;Accession: JC7104
R;Mulero, J.J.; Pace, A.M.; Nelken, S.T.; Loeb, D.B.; Correa, T.R.; Drmanac, Blochem. Blophys. Res. Commun. 263, 702-706, 1999
A;Title: ILLHY1: A novel interleukin-1 receptor antagonist gene.
A;Reference number: JC7104; MUID:99443727
A;Accession: JC7104
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A; Residues: 1-155 < MUL>
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61
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Pred. No. 2.3e-77;
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C;Keywords: alternative splicing; cytokine receptor; extracellular protei F;1-25/Domain: signal sequence #status predicted <SIG>
F;26-177/Product: interleukin-1 receptor antagonist #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Molecule type: protein
A;Residues: 26-52;70-77;122-127;170-175 <BIE>
A;Experimental source: culture medium, PMA-stimulated THP-1
C;Comment: For an alternative splice form, see PIR:A39386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: protein
A; Residues: 26-75; 97-108;110-116;120-131;163-176 <HAN>
R; Bienkowski, M.J.; Eessalu, T.E.; Berger, A.E.; Trueso
J. Biol. Chem. 265, 14505-14511, 1990
A; Title: Purification and characterization of interleul
A; Reference number: A37822; MUID:90354444
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A; Map position: 2914.2-2914.2
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A;Title: Interleukin-1 receptor antagonist A;Reference number: S08159; MUID:90136920 A;Accession: S08159
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A; Residues: 1-177 < LEN>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: mRNA
A; Residues: 1-177 <EI2>
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RESULT
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Best Local (
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                                                                  151
                                                                                                             126 EADQPVRLTQLPENGGWNAPITDFYFQQ 153
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                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                  CLSC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVP 125
                                                                                                                                                                                                                                                    FRIWDVNQKTFYLRNNQLVAGYLQGPNVNLEEKIDVVP-----
                                                                                                                                                                                                                                                                                          FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP--VILGVQGGSQ 66
                                                                                                                                                            CLSCVKSGDETRLQLEAVNITDLSENRKQDKRFAFIRSDSGPTTSFESAACPGWFLCTAM
                                                                  EADQPVSLTNMPDEG - - - VMVTKFYFQE
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0368; MUID:90220867
                                                                                                                                                                                                                                                                                                                                                               38.8%;
                                                                                                                                                                                                                                                                                                                                            13;
                                                                                                                                                                                                                                                                                                                                        Score 319.5;
Pred. No. 1.7e
13; Mismatches
                                                                  175
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                                                                                                                                                                                                                                                                                                                                                           1.7e-25;
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                                                                                                                                                                                                                                                  -IEPHALFLGIHGGKM 90
                                                                                                                                                                                                                                                                                                                                            Indels
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the mature protein, were con:
; Brewer, M.T.; Hannum, C.H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          receptor level
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Laborde, A.L.; Sl.
I.M.; Heinrikson,
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jon, R.L.;
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                                        A; Accession: B40956
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A;Map position: 2q14.2-2q14.2 (Superfamily: interleukin-1 (C;Kuperfamily: interleukin-1 (C;Kupwords: alternative splicing; cytokine receptor F;1-180/Product: interleukin-1 receptor antagonist, long F;1-3,25-180/Product: interleukin-1 receptor antagonist,
                                                                                                                                                                                                                                                                                                                     C;Date: 09-Sep-1994 #sequence_revision 09-Sep-1994 #text_change 16-Jul-1999 C;Accession: A44610; B40956; A49031; I56106; I52970 R;Matsushime, H.; Roussel, M.F.; Matsushima, K.; Hishinuma, A.; Sherr, C.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: mRNA
A; Residues: 1-3,25-180 < HAS>
A; Cross-references: GB: M55646; NID: g186291;
C; Comment: For an alternative splice form, s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Haskill, S.; Martin, G.; Van Le, L.; Morris, J.; Peace, A.; Bigler, C.F.; Jaffe, Proc. Natl. Acad. Sci. U.S.A. 88, 3681-3685, 1991
A;Title: cDNA cloning of an intracellular form of the human interleukin 1 receptor A;Reference number: A39386; MUID:91219436
A;Accession: A39386
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C;Species: Homo sapiens (man)
C;Date: 28-Feb-1992 #sequence_revision 11-Apr-1997 #text_change 26-May-2000
A;Cross-references: GB:M64404; NID:g198296; PIDN:AAA39277.1; R;Eisenberg, S.P.; Brewer, M.T.; Verderber, E.; Heimdal, P.; Proc. Natl. Acad. Sci. U.S.A. 88, 5232-5236, 1991
A;Title: Interleukin I receptor antagonist is a member of the A;Reference number: A40956; MUID:91271363
                                                                                                                                              A; Molecule type: mRNA
A; Residues: 1-178 <MAT>
                                                                                                                                                                                                                             A; TILLE: Cloning and expression of murine interleukin-1 A; Reference number: A44610; MUID:91316273
                                                                                                                                                                                                                                                                  Blood 78, 616-623, 19
A; Title: Cloning and
                                                                                                                                                                                                                                                                                                                                                                                                                C; Species: Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Gene: GDB: IL1RN
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A; Residues: 1-180 < RES>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Accession: I37893; A39386 R;Muzio, M.; Polentarutti,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Cross-references: GDB:125897;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       interleukin-1 receptor antagonist precursor
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Best Local S
Matches 73
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                                                                                                                                                                                                                                                                                                616-623, 1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FRIWDVNQKTFYLRNNQLVAGYLQGPNVNLEEKIDVVP-----IEPHALFLGIHGGKM 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP--VILGVQGGSQ 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                  names: IL-1Ra
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Pred. No. 1.7e-25;
3; Mismatches 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; PIDN:AAA59138.1; see PIR:A30368
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                                                                                        Brandhuber,
                                                                                                                   PID:g198297
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R; Zahedi, K.; Seldin, M.F.; Rits, M.; Ezekowitz, R.B.; Whitehead, A.S.
J. Immunol. 146, 4228-4233, 1991
A; Title: Mouse IL-1 receptor antagonist protein: Molecular characterization, A; Reference number: 156106; MUID:91250712
A; Accession: 156106
                                                                                                                             R;Eisenberg, S.P.; Brewer, M.T.; Verderber, E.; Heimdal, P.; Brandhuber, B.J.; Proc. Natl. Acad. Sci. U.S.A. 88, 5232-5236, 1991
A;Title: Interleukin 1 receptor antagonist is a member of the interleukin 1 gen A; Reference number: A40956; MUID:91271363
A;Accession: C40956
                                                                                                                                                                                                                                                                                                                                           RESULT
C40956
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C;Keywords: cytokine receptor
E;1-26/Domain: signal sequence #status predicted <SIG>
F;27-178/Product: interleukin-1 receptor antagonist #status predicted <MA2>
                     A;Cross-references: GB:M63101;
C;Superfamily: interleukin-1
                                                                                                                                                                                                                                               interleukin-1 receptor antagonist precursor
C;Species: Rattus norregicus (Norway rat)
C;Date: 20-Mar-1992 #sequence_revision 20-Ma
C;Accession: C40956
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Cytokine 6, 1-9, 1994
  C; Keywords:
                                                                  A; Molecule type: DNA
A; Residues: 1-178 <EIS>
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A; Residues: 1-178 <RE2>
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A; Accession: I52970
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: GB:S64082; NID:g238584; PIDN:AAB20265.1; PID:g238585 A;Experimental source: peritoneal macrophages, ICR strain A;Note: sequence extracted from NCBI backbone (NCBIN:64082, NCBIP:64085)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Title: Cloning, heterologous expression A;Reference number: A49031; MUID:92037824 A;Accession: A49031
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                                                                                                               A; Status: preliminary
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A; Residues: 23-178 <SHU>
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                   154 DRPVSLTNTPEE----PLIVTKFYFQE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       69
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DQPVRLTQLPENGGWNAP -- ITDFYFQQ 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SCAKSGDDIKLQLEEVNITDLSKNKEEDKRFTFIRSEKGPTTSFESAACPGWFLCTTLEA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FRIWDTNQKTFYLRNNQLIAGYLQGPNIKLEEKIDMVP----IDLHSVFLGIHGGKLCL 93
cytokine receptor
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                                            NID:g204928;
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Pred. No. 2.3e-24;
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                                                                                                                                                                                                                                                                    20-Mar-1992 #text_change 16-Jul-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                   176
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                                            PIDN: AAA41434.1;
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                                          PID:g204929
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interleukin-1 beta precursor - sheep
M;Alternate names: hematopoietin-1; IL-1 beta
C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Date: 08-Jun-1994 #sequence_revision 22-Nov-1996 #text_change 15-Oct-1999
C;Accession: S23010; S43047; S13092; B61246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   interleukin-1 receptor antagonist secreted form precursor - rabbit C;Species: Oryctolagus cuniculus (domestic rabbit) C;Date: 06-Oct-1994 #sequence_revision 18-Nov-1994 #text_change 16-Jul-1999 C;Accession: A54377; 146729 R;Cominelli, F:; Bortolami, M.; Pizarro, T.T.; Monsacchi, L.; Ferretti, M.; J. Biol. Chem. 269, 6962-6971, 1994 A;Title: Rabbit interleukin-1 receptor antagonist. Cloning, expression, A;Reference number: A54377; MUID:94165101 A;Accession: A54377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; Keywords: cytokine receptor; extracellular protein; F;1-25/Domain: signal sequence #status predicted <SIG> F;109/Binding site: carbohydrate (Asn) (covalent) #status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: mRNA
A; Residues: 1-177 <GOT>
A; Cross-references: GB:D21832; NID:g425787;
C; Superfamily: interleukin-1
C; Superfamily: interleukin-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Title: Interleukin-1 recetor antagonist in A;Reference number: I46729; MUID:93052512 A;Accession: I46729 A;Status: translated from GB/EMBL/DDBJ A;Status:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Note: sequence extracted from NCBI backbook; Goto, F.; Goto, K.; Miyata, T.; Ohkawara, Immunology 77, 235-244, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Molecule type: mRNA
A; Residues: 1-177 < COM>
A; Cross-references: GB: S68977; NID: g545740;
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Best Local S
Matches 67
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Best Local :
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                                                                                                                                                                                                                                EADQPVRLTQLPENGGWNAPITDFYFQQ 153
                                                                                                                                                                                                                                                                               CLSCVKSGDKMKLHLEAVNITDLGKNKEQDKRFTFIRSNSGPTTTFESASCPGWFLCTAL
                                                                                                                                                                                                                                                                                                                                                                                                      FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP--VILGVQGGSQ 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DHPVSLTNTPKE---PCTVTKFYFQE 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVPEA 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILGVQGGSQCL 68
                                                                                                                                                                                           EADQPVSLTNTPDD---SIVVTKFYFQE
                                                                                                                                                                                                                                                                                                                      CLSC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVP 125
                                                                                                                                                                                                                                                                                                                                                              FRIWDVNQKTFYLRNNQLVAGYLQGPNAKLEERIDVVP-----LEPQLLFLGIQRGKL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DOPVRLTQLPENGGWNAPITDFYFQQ 153
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Pred. No. 1e-
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C;Accession: JL0010; S01380
R;Maliszewski, C.R.; Baker, P.E.; Schoenborn, M.A.;
Mol. Immunol. 25, 429-437, 1988
A;Title: Cloning, sequence and expression of bovine
A;Reference number: A94695; MUID:88318652
A;Accession: JL0010
                                                                                                                                                                                                    interleukin-1 beta precursor - bovine
N;Alternate names: hematopoietin-1; IL-1 beta
C;Species: Bos primigenius taurus (cattle)
C;Date: 31-Mar-1989 #sequence_revision 31-Mar-1989 #text_change 22-Jun-1999
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A;Residues: 1-144,'L',146-266 <AND>
C;Comment: This protein lacks a conventional signal sequence for protein export. Cleavage ved form of interleukin-lbeta, unlike interleukin 1-alpha, is inactive.
C;Comment: Interleukin-lbeta precursor is less heavily myristoylated than interleukin-lacks.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: mRNA
A; Residues: 1-13, 'C', 15-54, 'K', 56-63, 'A', 65-144, 'L', 146-266 <
A; Residues: 1-13, 'C', 15-54, 'K', 56-63, 'A', 65-144, 'L', 146-266 <
A; Cross-references: EMBL:X54796; NID:g1273; PIDN:CAA38566.1;
R; Fiskerstrand, C.; Sargan, D.
Nucleic Acids Res. 18, 7165, 1990
A; Title: Nucleotide sequence of ovine interleukin-1 beta.
A; Reference number: S13092; MUID:g1088326
A; Accession: S13092
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A;Title: Nucleotide sequence of ovine macrophage interleukin-1 beta
A;Reference number: S23010; MUID:92119335
                      A; Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C; Keywords: cytokine; immunoregulation; inflammation; lymphokine; macrophage; mitogen F;114-266/Product: interleukin-1 beta *status predicted <MAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Title: Molecular cloning and characterization of ovine IL-lalpha and IL-lbeta A;Reference number: A61246; MUID:92120716
A;Accession: B61246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Note: the authors translated the codon AGT for residue 62 R;Andrews, A.E.; Barcham, G.J.; Brandon, M.R.; Nash, A.D. Immunology 74, 453-460, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Molecule type: mRNA
A;Residues: 1-13,'C',15-54,'K',56-61,'S',63,'A',65-144,'L',146-266
A;Cross-references: EMBL:X54796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          submitted to the EMBL Data Library, May 1992 A; Reference number: $43047
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A; Note: the sequence from Fig. 1
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A; Residues: 1-266 <SE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Seow, H.F.; Rothel, J.S.; David, M.J.; Wood, 
5NA Seg. 1, 423-426, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C; Superfamily: interleukin-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Reference number: A; Accession: S43047
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                              249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       190
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16 LKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILGVQGGSQCLSC-GVGQ 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 17.2%;
Local Similarity 31.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IL-1-beta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TPTLQLEEVD-PKVYPKRNMEKREVEYKTEIKNTVEFESVLYPNWYISTSQIEEKPVFLG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             QLPENGGWNAPITDF 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LKALHLPSQEMSREVVFCMSFVQGEERD-----
                                                                                                                                                                                                                                                                                                                                                                                                                              RF--RGGQD--ITDF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVPEADQPVRLT 134
1 - 266
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  AMAL>
                                                                                                                                                                                                                                                                                                                                                                                                                           259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NID:g1808; PIDN:CAA40293.1; PID:g1809 is inconsistent with that from Fig. 2 in having an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 141.5;
Pred. No. 5.7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ----NKIPVALGIRDKNLYLSCVKKGD 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .7e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             54;
                                                                                                   interleukin 1-alpha and interleukin
                                                                                                                                                     Davis, B.S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 266;
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                                                                                                                                                     Cosman, D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <FIS>
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                                                                                                                                                     Gillis,
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interleukin-1 beta precursor - pig
C:Species: Sus scrofa domestica (domestic pig)
C:Date: 20-May-1994 #sequence_revision 01-Dec-1995 #text_change 16-Jul-1999
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A;Residues: 1-251, 'A', 253-266 <LEO>
A;Cross-references: EMBL:X12498; NID:g448; PIDN:CAA31018.1; PID:g449
C;Comment: This protein is a cytokine that mediates a variety of immunoregulatory
C;Comment: This protein lacks a conventional signal sequence for protein export
ved form of interleukin-lbeta, unlike interleukin 1-alpha, is inactive.
C;Comment: Interleukin-lbeta precursor is less heavily myristoylated than interle
C;Superfamily: interleukin-1
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                                                                                                                                                                                                   Вb
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Eur. J. Biochem. 217, 45-52, 1993
A; Title: Gene sequence, cDNA construction,
A; Reference number: S38373; MUID:94039070
A; Accession: S38373
A; Status: preliminary
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Вb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: DNA
A; Residues: 1-267 < VAN>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Reference number: S01380; MUID:89016591 A; Accession: S01380
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A;Title: The nucleotide sequence for the cDNA of bovine interleukin-1 beta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: EMBL:X74568; NID:g407899;
                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                       Matches
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Best Local Similarity
235 YISTSQAEQKPVFL---GNSKGRQDITDF
                                               120 FLCTVPEADQPVRLTQLPENGGWNAPITDF 149
                                                                                               176 IKGKNLYLSCVMKDDTPTLQLEDVD-PKSYPKRDMEKREVEYKTEIKNRVEFESALYPNW
                                                                                                                                                                                                   132 LVLAG-----PHELKALHLLKGDLKREVVFCMSFVQGDD----
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                                                                                                                            61 VQGGSQCLSCGVGQE-PTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 119 ::| : | | | | | | | | | | | | | |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        75 EPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVPEADQPVRLT 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16 LKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILGVQGGSQCLSC-GVGQ 74
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                                                                                                                                                                                                                                                     MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILG 60
                                                                                                                                                                                                                                                                                                          44;
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                                                                                                                                                                                                                                                                                                     15.8%; Score 130; DB 2; ilarity 29.3%; Pred. No. 8.7e-06; Conservative 21; Mismatches 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                  Length 267
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R;Huether, M,J; Lin, G.; Smith, D.M.; Murtaugh, M.P.; Molitor, T.W.
Gene 129, 285-289, 1993
A;Title: Cloning, sequencing and regulation of an mRNA encoding porcine interleukin-1 be
A;Reference number: JN0724
A;Accession: JN0724
A;Accession: JN0724
A;Molecule type: mRNA
A;Residues: 1-267 <HUE>
A;Cross-references: GB:M86725; NID:g164607; PIDN:AAA02584.1; PID:g164608
A;Experimental source: alveolar macrophage
C;Comment: This protein is a pleiotropic cytokine that mediates a variety of processes if cytokine of interleukin-1beta, unlike interleukin 1-alpha, is inactive.
C;Comment: Interleukin-1beta, unlike interleukin 1-alpha, is inactive.
C;Comment: Interleukin-1beta precursor is less heavily myristoylated than interleukin-1a C;Reywords: cytokine; immunoregulation; inflammation; lipoprotein; lymphokine; macrophag
F;115-267/Product: interleukin-1 beta #status predicted <ILi>F;77/Binding site: myristate (Lys) (covalent) #status predicted
                                                                                                                                                   δÃ
                                                                                                                                                                                                                                                                                                                                           A;Cross-references: DDBJ:D42165; NID:g2463549; PIDN:BAA22528.1; PID:g2463550 C;Comment: This protein mediates a variety of physiological response to infections and synthesis by hepatocytes, and stimulation of chondrocytes and synovial cells to produce C;Superfamily: interleukin-1
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                                                  Q
                                                                                                       Вþ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R; Kato, H.; Youn, H.Y
Gene 177, 11-16, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Species: Equus caballus (domestic horse)
C;Date: 28-Oct-1997 #sequence_revision 28-Oct-1997 #text_change 20-Jun-2000
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C;Species: Sus scrofa domestica (domestic pig)
C;Date: 14-Jul-1994 #sequence_revision 22-Nov-1996 #text_change 22-Jun-1999
C;Accession: JN0724
                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: mRNA
A; Residues: 1-214 <KAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Reference number: A; Accession: JC5646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Title: Identification of an A; Reference number: JC5646; MC
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                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                       Query Match
Best Local Similarity
     158
                                                                                                          108
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                                                     96
                                                                                                                                                               37
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                                                                                                                                                   IKGEEISVVPNRWLDASLSPVILGVQGGSQCLSCGVGQ-EPTLTLEPVNIMELYLGAKES 95
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KRFVFNKMEIKGNVEFESAMYPNWYISTSQAEKSPVFLGN--TRGG--RDITDFIME 210
                                               KSFTFYRRDMGLTSSFESAAYPGWFLCTVPEADQPVRLTQLPENGGWNAPITDFYFQ 152
                                                                                                          VQGEE-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IKGKNLYLSCVMKDNTPTLQLEDID-PKRYPKRDMEKRFVFYKTEIKNRVEFESALYPNW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILG 60
                                                                                                                                                                                                                 39;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  H.Y.; Ohashi, T.; Watari, T.;
                                                                                                    ETDKIPVALGLKEKNLYLSCGMKDGKPTLQLETVD-PNTYPKRKME 157
                                                                                                                                                                                                                                       15.1%;
33.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15.7%;
28.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         an alternatively spliced transcript of equine interleukin-1 MUID:97080493
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                                                                                                                                                                                                                                    Score 124.5;
Pred. No. 2.4
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Pred. No. 1.1e-05;
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                                                                                                                                                                                                                                                                      DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Goitsuka, R.; Tsujimoto, H.; Hasegawa,
                                                                                                                                                                                                              48;
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A;Title: Nucleotide sequence of human monocyte interleukin 1 precursor cI A;Reference number: I51852
A;Recession: I51852
A;Scatus: translated from GB/EMBL/DDBJ
A;Recidus: translated from GB/EMBL/DDBJ
A;Residus: 1-5,'K',7-19,'H',21-110,'Q',112-176,'A',178-213,'P',215-269 < A;Residus: 1-5,'K',7-19,'H',215-269 < A;Residus: 1-5,'K',7-19,'H',215-269
A;Cross-references: EMBL:X56087; NID:g35662; PIDN:CAA39567.1; PID:g35663 R;Zsebo, K.M.; Wypych, J.; Yuschenkoff, V.N.; Lu, H.; Hunt, P.; Dukes, PBlood 71, 962-968, 1988
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A; Residues: 1-5, K, 7-269 < AUR>
A; Residues: 1-5, K, 7-269 < NID: 9186268; PIDN: AAA36106.1; PID: 9307043
A; Cross-references: GB: K02770; NID: 9186268; PIDN: AAA36106.1; PID: 9307043
R; March, C.J.; Mosley, B.; Larsen, A.; Cerretti, D.P.; Braedt, G.; Price,
Nature 315, 641-647, 1985
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A; Title: Human interleukin-1 beta gene.
A; Reference number: A29019; MUID:87248099
A; Accession: A29019
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C;Date: 28-Feb-1986 #sequence_revision 15-May-1998 #text_change 15-Sep-2000
C;Date: 28-Feb-1986 #sequence_revision 15-May-1998 #text_change 15-Sep-2000
C;Accession: A25542; A29019; A94023; A93361; I51852; I65200; I38132; B27616;
C;Accession: A25542; A29019; A94023; A93361; I51852; I65200; I38132; B27616;
R;Clark, B.D.; Collins, K.L.; Gandy, M.S.; Webb, A.C.; Auron, P.E.
Nucleic Acids Res. 14, 7897-7914, 1986
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A;Molecule type: mRNA
A;Residues: 1-269 <NIS>
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A; Title: Nucleotide sequence of human monocyte interleukin 1 precursor cDNA
A; Reference number: A94023; MUID:85088517
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Proc. Natl. Acad. Sci. U.S.A. 81, 7907-7911, 1984
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A;Note: the mRNA sequence had codon AAG for
R;Bensi, G.; Raugei, G.; Palla, E.; Carinci,
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A;Accession: I38132
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R; Kotenko, S.V.; Bulenkov ii, S.A.; Vinetskii, Y.P.
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A; Residues: 1-269 <MAR>
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A; Residues: 1-269 <BEN>
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A; Residues: 1-5, 'K', 7-269 < CLA>
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N;Alternate names: hematopoietin-1; IL-1 be
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A;Map position: 2q13-2q21
A;Introns: 16/2; 33/3; 101/1; 156/1; 199/3
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A;Title: The 31-KDa precursor of interleukin lalpha is myristoylated on specific lysines A;Reference number: A48293; MUID:93348250
A;Contents: annotation; myristylation of lysines R;Nanduri, V.B.; Hulmes, J.D.; Pan, Y.C.E.; Kilian, P.L.; Stern, A.S.
Biochim. Biophys. Acta 1118, 25-35, 1991
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J. Biol. Chem. 266, 7081-7086, 1991
A;Title: Structure-function mapping of interleukin 1 precursors.
A; Reference number: A39774; MUID:91201363
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                                                                         RESULT
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A; Residues: 117-123, 'X', 125-126, 'X', 128 <ZSE>
interleukin-1 beta precursor
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29.0%;
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Pred. No. 4.6e-05;
3; Mismatches 61
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R;Gray, P.W.; Glaister, D.; Chen, E.; Goeddel, D.V.; Pennica, J. Immunol. 137, 3644-3648, 1986
A;Title: Two interleukin 1 genes in the mouse: Cloning and ex. A;Reference number: 155969; MUID:87058957
A;Accession: 155969
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A. Molecule type: mRNA
A. Rolecule type: mRNA
A.; Residues: 1-268 < YOU>
C.; Comment: This protein lacks a conventional signal sequence for protein export.
ved form of interleukin-lbeta, unlike interleukin 1-alpha, is inactive.
C.; Comment: Interleukin-lbeta precursor is less heavily myristoylated than interle
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A;Title: Rabbit II-1. Cloning, expression, biologic properties, A;Reference number: A30584; MUID:89176242
A;Accession: A30584
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                                                                                                                                                                                                                                                                                                                                                                                       C;Species: Mus musculus (house mouse)
C;Date: 26-Jul-1996 #sequence_revision 22-Nov-1996 #text_change
C;Accession: 155969; A24719; S13029
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Protein Eng. 2, 545-551, 1989
A;Title: Cloning of rabbit interleukin-1 beta:
A;Reference number: A94230; MUID:89315718
A;Accession: JU0082
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C;Date: 25-May-1989 #sequence_revision 22-Nov-1996 #text_change
C;Accession: A27714; A30584; JU0082; A32166
A; Reference number: A24719; MUID:87117546
A; Accession: A24719
                                                                             R;Telford, J.L.; Macchia, G.; Massone, Nucleic Acids Res. 14, 9955-9963, 1986
                                                                                                                                                                   A; Molecule type: mRNĀ
A; Residues: 1-269 < RES>
                                                                                                                                                                                                                            A; Status: preliminary;
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N;Alternate names: hematopoietin-1; IL-1 beta
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A; Cross-references: GB: M26295; NID: g516632;
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A; Residues: 1-268 < MOR>
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                                                  A; Title: The murine interleukin 1-beta gene: structure and
                                                                                                                                   A; Cross-references: GB:M15131; NID:g198293; PIDN:AAA39276.1; PID:g309398
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Pred. No. 9.3e-05;
1; Mismatches 63;
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complement C3-Q2 - common carp C;Species: Cyprinus carpio (common carp) C;Date: 22-Oct-1999 #sequence_revision 22-Oct-1999 #text_change 22-Oct-1999 C;Accession: T30517 R;Nakao, M.; Obo, R.; Mutsuro, J.; Fujiki, K.; Nonaka, M.; Yano, T. submitted to the EMBL Data Library, July 1998 A;Description: Sequence diversity of the third complement component (C3) from a bony fis 밁 . δ 밁 δÃ 밁 Qy A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-1230 <NAK>
A;Cross-references: EMBL:AB016215; NID:d1253045; PID:d1037610; PIDN:BAA36623.1 ğ Ъ Ş A;Gene: IL-1-beta
C;Superfamily: interleukin-1
C;Keywords: cytokine; immunoregulation; inflammation; lymphokine; macrophage; mitogen
F;118-269/Product: interleukin-1 beta #status experimental <IL1> A;Status: preliminary
A;Molecule type: protein
A;Rolecule type: protein
A;Residues: 118-269 < CAU>
A;Residues: 118-269 < CAU>
C:Comment: This protein lacks a conventional signal sequence for protein export. Cleavag ved form of interleukin-lbeta, unlike interleukin 1-alpha, is inactive.
C:Comment: Interleukin-lbeta precursor is less heavily myristoylated than interleukin-lackin-A;Molecule type: mRNA A;Residues: 1-269 <TEL> A;Cross-references: GB:X04964; NID:g52666; PIDN:CAA28637.1; PID:g52667 R;Daumy, G.O.; Wilder, C.L.; Merenda, J.M.; McColl, A.S.; Geoghegan, K.F.; Otterness, FEBS Lett. 278, 98-102, 1991 A; Reference number: Z20840 A; Accession: T30517 밁 Ъ QY A;Title: Reduction of biological activity of murine recombinant interleukin-lbeta by se. A;Reference number: S13029; MUID:91130610
A;Accession: S13029 C; Genetics: Query Match Best Local S Matches 48 Query Match 9.8%; Score 80.5; DB 2; Best Local Similarity 27.3%; Pred. No. 7.3; Matches 35; Conservative 17; Mismatches 53 673 789 118 GWFLCTVP 125 733 ASLHESIRKAVSFLEGRLPQLINPYAVAMMSYAMANENK----LNKDILMKHSSQHEAGR 788 233 EFPNWYISTSQAEHKPV---FLGNNSGQD--IIDF 262 115 174 PVALGLKGKNLYLSCVMKDGTPTLQLESVDPKQ-YPKKKMEKRFVFNKIEVKSKVEFESA 232 123 LHYRLRDEQQKSLVLSDPYELK-ALH----LNGQNI----NQQVIFSMSFVQGEPSNDKI 173 73 14 SALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP-VILGVQGGSQCLSCGV 72 56 Local Similarity 7 LCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLS------ 55 SW---TVP 793 SALKWLILHKQTLYGSFKEDSAVIQGEMVGGIQSKDADASLTAFVVIAMQEAREICAGSV 732 AYPGWFLCTVPEADQPVRLTQLPENGGWNAPITDF 149 PVILGVQGGSQCLSCGVGQ-EPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESA 114 Conservative GOEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYP 117 14.5%; Score 119.5; DB 1; Length 269; 31.0%; Pred. No. 0.00011; 23; Mismatches 53; Length 1230; Indels 23; 27; Gaps Gaps 4. 8

Search completed: June 25, 2001, 14:05:59
Job time: 61 sec

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CYSD_PSEAE
YTYA_TREHY
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P2Y7_HUMAN
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8 cercocebus
2 moraxella c
6 riftia pach
9 chlamydia p
6 methanosarc
8 plasmodium
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8 bos taurus
6 equus cabal
2 capra hircu
9 sus scrofa
0 macaca mula
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Schreuder H., Tardif C., Trump-Kallmeyer S., Soffientini A., Sarubbi E., Akeson A., Bowlin T., Yanofsky S., Barrett R.W.: "A new cytokine-receptor binding mode revealed by the crystal structure of the IL-1 receptor with an anti---
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Bienkowski M.J., Eessalu T.E., Berger A.E., Truesd Shelly J.A., Laborde A.L., Zurcher-Weely H.A., Rea Heinrikson R.L., Chosay J.G., Tracey D.E.;
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MEDILINE-92037824; PubMed=1834470;

Shuck M.E., Eessalu T.E., Tracey D.E., Bienkowski M.J.;

"Cloning, heterologous expression and characterization of murine interleukin 1 receptor antagonist protein.";

Eur. J. Immunol. 21:2775-2780(1991).
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MEDLINE=91250712; PubMed=1828262;

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Zahedi K., Seldin M.F., Rits M., Ezekowitz R.A., Whitehead A.S.;

Zahedi K., Seldin M.F., Rits M., Ezekowitz R.A., Whitehead A.S.;

"Mouse IL-1 receptor antagonist protein. Molecular characterization,

gene mapping, and expression of mRNA in vitro and in vivo.";

J. Immunol. 146:4228-4233(1991).
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MEDLINE=91316273; PubMed=1830498;

Matsushime H., Roussel M.F., Matsushima K., Hishinuma A.,

"Cloning and expression of murine interleukin-1 receptor
"Cloning and expression of murine interleukin-1 receptor
in macrophages stimulated by colony-stimulating factor 1.
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01-NOV-1997
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                                                                                                                                                                                                                                                                                                                                                                                                FUNCTION:
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bkine 6:1-9(1994).
                                                                         ; P18510; 1IRA.
MGI:96547; Il1
                                                                                                                                                                                                                                                                                                                       SWISS-PROT entry is copyright. It is produced sen the Swiss Institute of Bioinformatics and
                                                                                                                                                                                                                                                                                            non-profit institutions as long
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Rodentia;
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Last annotation updat
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Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                      Bioinformatics
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Matches 71
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029056;
15-DEC-1998 (Rel. 37, Cre
15-DEC-1998 (Rel. 38, La;
15-JUL-1999 (Rel. 38, La;
    DISULFID
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SEQUENCE
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CHAIN
                                                                                                                                                                                                                                                            entities requires a license agreement (Su or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                 the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                  This
                                                                                                                                                                                                                                                                                                                                                                                                                                        "Characterization of IRAP in morphine treated pig.";
Submitted (MAY-1996) to the EMBL/GenBank/DDBJ databases
-I-FUNCTION: IL-1RA INHIBITS THE ACTIVITY OF IL-1 BY B:
RECEPTOR. IL-1RA HAS NO IL-1 LIKE ACTIVITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sus scrofa (Pig).
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                       Glycoprotein;
SIGNAL
                                                                                                                                                                                                                           EMBL; L38849; AAA99424.1;
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                                                                                                                                          PROSITE; PS00253; INTERLEUKIN_1;
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37, Last sequence update)
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; 84AA002A3119C024 CRC64;
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No. 6.2
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      CRC64;
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Query Match Best Local Similarity

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
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15-JUL-1999 (Rel. 38, Last annotation update)
INTERLEUKIN-1 RECEPTOR ANTAGONIST PROTEIN PRECURSOR (IL-1RA) (IL-1RN)
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FUNCTION: IL-1RA INHIBITS THE ACTIVITY OF IL-1 BY BINDING TO RECEPTOR. IL-1RAHAS NO IL-1 LIKE ACTIVITY.
SIMILARITY: BELONGS TO THE IL-1 FAMILY.
CLSC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTYP 125
                                                                                FRMKDSALKYLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP--VILGVQGGSQ 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EADQPVRLTQLPENGGWNAPITDFYFQQ 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLSCVKSGDEMKLQLDAVNITDLRKNSEQDKRFTFIRSDSGPTTSFESAACPGWFLCTAL 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLSC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVP 125
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EADQPVGLTNTPKAA---VKVTKFYFQQ
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                                                   FRIWDVNQKIFYLRNNQLVAGYLQGPNTKLEEKIDVVP-----
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                                                                                                                                                                               Score 295.5;
Pred. No. 1.
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N-LINKED (GLCNAC. .
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ches 51;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Glycoprotein; Signal. SIGNAL 1 2
                                                                                                                                                                                                                                                                                                                                                                                                                                        CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; M63101; AAA41434.1;
PIR; C40956; C40956.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene family: evolution of a cytokine control mechanism.";
Proc. Natl. Acad. Sci. U.S.A. 88:5232-5236(1991).
-i-FUNCTION: IL-IRA INHIBITS THE ACTIVITY OF IL-1 BY BIN
RECEPTOR. IL-IRA HAS NO IL-1 LIKE ACTIVITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Brandhuber B.J., Thompson R.C.; Interleukin 1 receptor antagonist is a member of the interleukin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mEDLINE=91271363; PubMed=1828896; Eisenberg S.P., Brewer M.T., Verd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rattus norvegicus (Rat).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ol-MAY-1992 (Rel. 22, Created)
Ol-MAY-1992 (Rel. 22, Last sequence update)
Ol-MOV-1995 (Rel. 32, Last annotation update)
INTERLEUKIN-1 RECEPTOR ANTAGONIST PROTEIN PRECURSOR (IL-1RA) (IL-1RN)
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PROSITE; PS00253; INTERLEUKIN_1; 1.
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Pfam; PF00340; inter
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                                                                                                                                                                                   FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILGVQGGSQCL
DHPVSLTNTPKE - - - PCTVTKFYFQE
                                      DQPVRLTQLPENGGWNAPITDFYFQQ 153
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                                                                                                    SC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVPEA
                                                                                                                                                                FRIWDTNQKTFYLRNNQLIAGYLQGPNTKLEEKIDMVP----IDFRNVFLGIHGGKLCL
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                                                                                 SCVKSGDDTKLQLEEVNITDLNKNKEEDKRFTFIRSETGPTTSFESLACPGWFLCTTLEA
                                                                                                                                                                                                                                                  67; Conserv
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178 #
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Pred. No. 2.8e-23;
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F3A5754FB6C51B03 CRC64;
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01-AUG-1992 (Rel. 23, Created)
01-AUG-1992 (Rel. 23, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
INTERLEUKIN-1 RECEPTOR ANTAGONIST PROTEIN PRE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; S68977; AAB30093.1;
EMBL; M57526; AAA31374.1;
EMBL; D21832; BAA04860.1;
PIR; A54377; A54377.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=93052512; PubMed=1427977; Goto F., Goto K., Miyata T., Ohkawara S., Takao T., Mori S., Furukawa S., Madda T., Twanaga S., Shimonishi Y., Yoshinaga M.; "Interleukin-1 receptor antagonist in inflammatory exudate cells rabbits. Production, purification and determination of primary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Elbetween the Bioinformatics and the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A. Hamada H., Mulligan | Submitted (XXX-1992)
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Brewer M.T., Eisenberg S.P., Ng R.K.;
"Rabbit interleukin-1 receptor antago
                                                                                                                                                                                                                                                                                                                                                                                        Glycoprotein;
SIGNAL
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Eukaryota; Metazoa; Chordáta; Craniata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the European Bioinformatics Institute.
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FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP--VILGVQGGSQ 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PF00340;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     PR00264; INTERLEUKIN1.
; PS00253; INTERLEUKIN_1; 1.
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109
177
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                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                          ; Signal.
1 25
26 177
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AAA31374.1; -.
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109 I
20214 MW;
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                                                               17;
                                                                                                                                                                                                                PROTEIN.
BY SIMILARITY.
N-LINKED (GLCNAC. .
W; F5BC087F097FFEAF C
                                                               Score 286.5;
Pred. No. 1.2e
7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                        BY SIMILARITY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     There are no rest
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vertebrata; Eu
ae; Oryctolagus
                                                                                          ; DB 1
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Query Match
Best Local S
Matches 65
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018999; 077745;
15-DEC-1998 (Rel. 3
15-DEC-1998 (Rel. 3
15-JUL-1999 (Rel. 3
                                                                 DISULFID
CARBOHYD
CONFLICT
SEQUENCE
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SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
MEDLINE-97366446; PubMed-9223227;
Kato H., Ohashi T., Matsushiro H.,
Haseqawa A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     _HORSE
                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. There are no restructed by non-profit institutions as long as its content modified and this statement is not removed. Usage by and
                                                                                                                                                                                                                                                                                                                                                                                Howard R.D., McIwraith C.W., Trotter G.W., Nyborg J.K.;
"Cloning of equine interleukin-1 receptor antagonist and
determination of its full-length cDNA sequence.";
Am. J. Vet. Res. 59:712-716(1998).
-i- FUNCTION: IL-1RA INHIBITS THE ACTIVITY OF IL-1 BY BII
RECEPTOR. IL-1RA HAS NO IL-1 LIKE ACTIVITY.
-i- SIMILARITY: BELONGS TO THE IL-1 FAMILY.
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Mammalia; Eutheria;
NCBI_TaxID=9796;
                                                                                                                                      CHAIN
                                                                                                                                                    Glycoprotein; SIGNAL
                                                                                                                                                                                                                                EMBL; U83714; BAA22529.1; -. EMBL; U92482; AAC39257.1; -.
                                                                                                                                                                                                                                                                      entities requires a license agreement (Some or send an email to license@isb-sib.ch).
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15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1999 (Rel. 38, Last annotation update)
INTERLEUKIN-1 RECEPTOR ANTAGONIST PROTEIN PRECURSOR (IL-1RA) (IL-1RN)
                                                                                                                                                                            PRINTS; PR00264; INTERLEUKIN1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
                                                                                                                                                                                                      Pfam; PF00340;
                                                                                                                                                                                                                      InterPro; IPR000975;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tsujimoto H., Hasegawa A.;
"Molecular cloning and functional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=98285942; PubMed=9622739;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      receptor antagonist."
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109
19
                                                                                                                                                               Signal
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interleukin1
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109
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            33.2%;
                                                                   20459
                                                                   MW.
                                                              N-LINKED (GLCNAC. .
F -> L (IN REF. 2).
; 1ABC377F1F1CF80B
           Score 273.5;
Pred. No. 2.6
                                                                                                         PROTEIN.
BY SIMILARITY
                                                                                                                                    BY SIMILARITY.
INTERLEUKIN-1 RECEPTOR ANTAGONIST
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65;

Conservative

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Similarity

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                                                                                                                                       EMBL; X54796; CAA38566.1; -
EMBL; X56972; CAA42293.1; -
PIR; S13092; S13092.
PIR; S13810; S13810.
PIR; S23010; S23010.
HSSP; P01584; 411B.
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between
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P21621;
01-MAY-1991 (Rel. 18, Created)
01-MAR-1992 (Rel. 21, Last sequence updat
30-MAY-2000 (Rel. 39, Last annotation upc
INTERLEUKIN-1 BETA PRECURSOR (IL-1 BETA)
                                                                                                                                                                                                                                                                                                                                                                                                     the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Nucleotide sequence of ovine interleukin-1 beta.";
"Nucleic Acids Res. 18:7165-7165(1990),
"I FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES
THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL
MATURATION & PROLIFERATION BY INDUCING IL-1 REPORTED AS ENDOGENOUS PROGENS, AND ARE REPORTED TO STIMULATE
IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING
IDENTIFIED AS ENDOGENOUS PYROGENS, AND ARE REPORTED TO STIMULATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SHEEP
PRINTS;
                                                   InterPro; IPR000975; -.
InterPro; IPR002348; -.
Pfam; PF00340; interleu
                                                                                                                                                                                                                                                                                                                                               entities requires a license agreement (See or send an email to license@isb-sib.ch).
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[2]
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Seow H.F., Rothel J.S., David M.J., Wood P.R.;
"Nucleotide sequence of ovine macrophage interleukin-1 beta cDNA.";
DNA Seq. 1:423-426(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
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Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Caprinae; Ovis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ovis aries (Sheep).
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INTERLEUKIN-1 BETA PRECURSOR (IL-1 BETA).
                                                            modified and this statement is not removed. entities requires a license agreement (See or send an email to license@isb-sib.ch).
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                          mitted (MAR-1995) to the EMBL/GenBank/DDBJ databases. EUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STINULATES THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL MATURATION & PROLIFERATION, & FIBROBLAST GROWTH FACTOR ACTIVITY. IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING IDENTIFIED AS ENDOGENOUS PYROGENS, AND ARE REPORTED TO STIMULATE
                                                                                                                    European
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Cloning, sequence and expression of bovine interleukin 1 beta complementary DNAs."; Mol. Immunol. 25:429-437(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic Acids [2]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-MAR-1989
30-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gillis S., Cerretti D.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=88318652; PubMed=3261832; Maliszewski C.R., Baker P.E., Sch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bovidae; Bovinae;
NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ol-MAR-1989 (Rel. 10, Created)
01-MAR-1989 (Rel. 10, Last seq
30-MAY-2000 (Rel. 39, Last ann
INTERLEUKIN-1 BETA PRECURSOR (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
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                                                                                                                                                                                                                   SUBUNIT: MONOMER.

SUBUNIT: MONOMER.

DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.

MISCELLANEOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGMENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS
                                                                                                                                           SECRETORY PROTEINS.
SIMILARITY: BELONGS
                                                                                                                                                                                                                                                                                                                                                                                                                                       FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL MATURATION & PROLIFERATION, & FIBROBLAST GROWTH FACTOR ACTIVITY. IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING IDENTIFIED AS ENDOGENOUS PYROGENS, AND ARE REPORTED TO STIMULATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FUNCTION:
                                                                                                                                                                                                                                                                                                                                                                                                            THE RELEASE OF PROSTAGLANDIN AND
                                  SWISS-PROT entry is copyright. It is produced through a centre the Swiss Institute of Bioinformatics and the EMBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HF--RGGQD--ITDF 259
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             Bioinformatics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chordata; Craniata; Vertebrata; Euteleostomi;
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31.9%;
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Last annotation update)
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Pred. No. 2.6e
22; Mismatches
             Institute.
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                                                                                                                                                                                                                DIFFERING
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BY DAMAGED CELLS
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                                                                  a collaboration
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tions on its
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Best Local
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PRINTS; PRO0264; INTERLEUKIN1.; 1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
Cytokine; Macrophage; Mitogen; Inflat
PROPEP 1 113
CHAIN 114 266
CONFLICT 252 252
A -> G
SEQUENCE 266 AA; 30774 MW; 9D1EF
        MEDLINE-97080493; PubMed-8921838; Kato H., Youn H.Y.. Obert
                                                                                                                                                                                                                                                                        IL1B_HORSE STANDARD; PRT; 268 AA  
Q28386; 077744; 018995;  
15-DEC-1998 (Rel. 37, Created)  
15-DEC-1998 (Rel. 37, Last sequence update)  
30-MAY-2000 (Rel. 39, Last annotation updat-INTERLEUKIN-1 BETA PRECURSOR (IL-1 BETA).
                                                       Howard R.D., McIlwraith C.W., Trotter G
"Cloning of equine interleukin-1 alpha
and determination of their full-length
and J. Vet. Res. 59:704-711(1998).
                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PIR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 use by modified
                                                                                                                                                Tsujimoto H., Hasegawa A.;
"Molecular cloning of equi
                                                                                                                                                                                                                                                    Equus caballus (Horse).
                                                                                                                                                                                 SEQUENCE FROM N.A.
MEDLINE=96131982; PubMed=8578682;
                                                                                                                                                                                                                   NCBI_TaxID=9796;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pfam; PF00340;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entities requires a
                                                                                                    MEDLINE=98285941; PubMed=9622738;
                                                                                                             SEQUENCE FROM N.A.
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S01380; S01380.
; P01584; 1HIB.
                                                                                                                                                                        H., Ohashi T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LKYLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILGVQGGSQCLSC-GVGQ
                                                                                                                                       Immunol.
                                                                                                                                                                                                                                                                                                                                                                                                               QLPENGGWNA - - PITDF
                                                                                                                                                                                                                                                                                                                                                                                                                                    TPTLQLEEVD-PKVYPKRNMEKRFVFYKTEIKNTVEFESVLYPNWYISTSQIEERPVFL-
                                                                                                                                                                                                                                                                                                                                                                                                                                                         EPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVPEADQPVRLT 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LKALHLLSQEMNREVVFCMSFVQGEERD-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           s requires a license agreement (S an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                          ---GHFRAGQDITDF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                non-profit institutions as long and this statement is not removed
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                                                                                                                                     Immunopathol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             interleukin-1;
                                                                                                                                                                                                                           Chordata; Craniata; Vertebrata; Euteleostomi; Perissodactyla; Equidae; Equus.
                                                                                                                                                                      Nakamura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16.5%;
                                                                                                                                                                                                                                                                                                                                                                                                             149
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                                                                                                                                     interleukin-1 a
48:221-231(1995)
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Pred.
                                                                                                                                                                      Ν.,
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A -> G (IN REF. 2).
9D1EF8F575070586
                                                                                         Trotter G.W.,
              Watari
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Inflammatory
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                                                                                                                                                                      Nishimura Y.,
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No. 8.
            Τ.,
                                                                                                                                                                                                                                                                                     update)
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                                                                    cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .5e-07
            Goitsuka
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                                                                             equine interleukin-1
                                                                                                                                              alpha
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                                                                                        Nyborg J.K.;
                                                                    sequences
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                                                                                                                                                                     Watari T.,
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                                                                                                                                                                      Goitsuka R.,
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PRINTS; PR00264; INTERLEUKIN1.
PROSITE; PS00253; INTERLEUKIN_1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entities requires a license agreement (See http://www.isb-sib.ch/announce, or send an email to license@isb-sib.ch).
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Gene 177:11-16(1996)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Identification of an alternatively spliced transcript of equine interleukin-1 beta.";
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MISCRILANGOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGMENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER
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                                                                                                 FLCTVPEADQPVRLTQLPENGGWNAPITDFYFQ 152
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                                                                                                                            LKEKNLYLSCGMKDGKPTLQLETVD-PNTYPKRKMEKRFVFNKMEIKGNVEFESAMYPNW
                                                                                                                                                        VQGGSQCLSCGVGQ-EPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 119
                                                                                                                                                                                    LVLSGA-----CELQAVHLNGENTNQQVVFCMSFVQGEE-----
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D -> N (IN REF. 2).
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AM -> VV (IN REF. 2).
V -> M (IN REF. 2).
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EG -> V (IN REF. 2).
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M -> V (IN REF. 2).
S -> K (IN REF. 2).
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Pred. No. 2e-06;
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INTERLEUKIN-1 BETA.
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15-DEC-1998 (Rel. 3
30-MAY-2000 (Rel. 3
INTERLEUKIN-1 BETA
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SEQUENCE
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Mammalia; Eutheria;
Bovidae; Caprinae; (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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PROSITE; PS00253; INTERLEUKIN_1;
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DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENDS OF THESE PROFEINS SERVE SOME AS YET UNDEFINED FUNCTION.

MISCELLANEOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGMENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER SECRETORY PROTEINS.
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                                         QLPENGGWNAPITDF
                                                                                  TPTLQLEEVD-PKVYPKRNMEKRFVFYKTEIKNTVEFESVLYPNWYISTSQIEEKPVFLG
                                                                                                                                                                                                         LKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILGVQGGSQCLS-CGVGQ 74
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HF -- RGGQD -- ITDF
                                                                                                                           EPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVPEADQPVRLT 134
                                                                                                                                                                    LKALHLLSQEMSREVVFCMSFVQGEERD------NKIPVALGIRDKNLYLSWVKKGD 189
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42; Conserv
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30769 MW;
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31.1%;
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01-AUG-1992
01-AUG-1992
30-MAY-2000
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-1- FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES
-1- FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, B-CELL
THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL
MATURATION & PROLIFERATION, & FIBROBLAST GROWTH FACTOR ACTIVITY.

IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING
ILDENTIFIED AS ENDOGENOUS PYROGENS, AND ARE REPORTED TO STIMULATE
THE RELEASE OF PROSTAGLANDIN AND COLLAGENASE FROM SYNOVIAL CELLS.
                                                                                                                                                                                                                                                                                 Cytokine; Macrophage; Mitogen; Inflammatory response; PROPEP 1 114 BY SIMILARITY.
CHAIN 115 267 INTERLEUKIN-1 BETA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the European Bioinformatics Institute. There are no resuse by non-profit institutions as long as its content modified and this statement is not removed. Usage by a
                                                                                                                                                                                                                                                                                                                             PRINTS; PR00262; IL1HBGF.
PRINTS; PR00264; INTERLEUKIN1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
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PIR; JN0724; JN0724.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nuecner M.J., Lin G., Smith D.M., Murtaugh M.P., Molitor T.W., "Cloning, sequencing and regulation of an mRNA encoding porcine interleukin-1 beta.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A. MEDLINE=93314975; PubMed=8325511;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota;
Mammalia;
                                                                                                                                                                                                                                                                     SEQUENCE
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DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.

MISCELLANEOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SECHENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER SECRETORY PROTEINS.
YISTSQAEQKPVFL----
                                                                                                                          LVLAG-----PHMLKALHLLTGDLKREVVFCMSFVQGDDSN-----NKIPVTLG
                                                                                                                                                       MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILG 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SWISS-PROT entry is copyright. It is produced through a collaboration -
                              FLCTVPEADQPVRLTQLPENGGWNAPITDF
                                                               IKGKNLYLSCVMKDNTPTLQLEDID-PKRYPKRDMEKRFVFYKTEIKNRVEFESALYPNW
                                                                                           VQGGSQCLSCGV-GQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 119
                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eutheria;
                                                                                                                                                                                                                                                                    115
267
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (Rel. 23, Created)
(Rel. 23, Last sequence up)
(Rel. 39, Last annotation
                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                    ΑA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23, Created)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chordata; Craniata; Vertebrata; Euteleostomi; Cetartiodactyla; Suina; Suidae; Sus.
                                                                                                                                                                                                                                                                    30404 MW;
                                                                                                                                                                                                      15
28
                                                                                                                                                                                                      3.78;
                                                                                                                                                                                       23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                 INTERLEUKIN-1 BETA.
; 7F6B92B784D5086F CRC64;
                                                                                                                                                                                                      Score 129;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT;
                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                    Gaps
                                                             234
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                                                                                                                                            Matches
                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-FEB-1996
01-FEB-1996
15-DEC-1998
                                                                                                                                                                                                                  CHAIN
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IL1B_MACMU P48090;
                                                                                                                                                                                                                                                                                                                                                                                                                   use by non-profit institutions as long modified and this statement is not removed entities requires a license agreement (See
                                                                                                                                                                                                                                                                                                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entre European Bioinformatics Institute. There are no resti
                                                                                                                                                                                                                                                                       PRINTS; PR00262; IL1HBGF.
PRINTS; PR00264; INTERLEUKIN1
PROSITE; PS00253; INTERLEUKIN
                                                                                                                                                                                                                                                                                                                                                            EMBL; U19845; AAA86709.1; HSSP; P01584; 1HIB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ÷
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-FEB-1996 (Rel. 33, Last sequence update) 15-DEC-1998 (Rel. 37, Last annotation updat INTERLEUKIN-1 BETA PRECURSOR (IL-1 BETA).
                                                                                                                                                                                                                                                                                                                    Pfam; PF00340;
                                                                                                                                                                                                                                                                                                                                                                                                      or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       J. Immunol. 155:3946-3954(1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nonhuman primates."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=96003435; PubMed=7561102;
Villinger F.J., Brar S.S., Mayne A.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE=Blood;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                              PROPEP
                                                                                                                                                                                                                                                           Cytokine;
                                                                                                                                                                                                                                                                                                                                InterPro;
                                                                                                                                                                                                                                                                                                                                               InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Comparative sequence analysis of cytokine genes from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9544;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cercopithecinae; Macaca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Macaca mulatta (Rhesus macaque).
  120
                               178
                                                                                    134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IL/1B
                                                        61
                                                                                                   1 MYLSGALCFRMKDSALKYLYLHNNQLLAGGLHAGKVIKGEEISVYPNRWLDASLSPYILG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SECRETORY PROTEINS.
SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.

MISCELLANGOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGMENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS
OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES THYMOCITE PROLIFERATION BY INDUCING IL-2 RELASE, B-CELL MATURATION & PROLIFERATION, & FIBROBLAST GROWTH FACTOR ACTIVITY. IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING IDENTIFIED AS ENDOGENOUS PYROGENS, AND ARE REPORTED TO STIMULATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SUBUNIT: MONOMER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    THE RELEASE OF PROSTAGLANDIN AND COLLAGENASE
FLCTVPEADQPVRLTQLPENGGWNA--PITDFYFQ
                            LKAKNLYLSCVLKDDKPTLQLESVD-PKNYPKKKMEKRFVFNKIEINNKLEFESAQFPNW
                                                       VQGGSQCLSCGV-GQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGW 119
                                                                                                                                                                                                                                                                                                                                 IPR000975; -. IPR002348; -.
                                                                                                                                                                                                                                            Macrophage; Mitogen;
                                                                                                                                                                                                                  117
269
                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (Rel. 33, Created)
                                                                                                                                                                                                                                                                                                                 interleukin-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STANDARD;
                                                                                                                                                                                                                                                                       INTERLEUKIN_1; 1
                                                                                   ELKALHLQGQDLEQQVVFSMSFVQGEESN
                                                                                                                                                                                                                                  269
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chordata;
Primates;
                                                                                                                                                                                                                  30481 MW;
                                                                                                                                                      15.4%;
29.7%;
                                                                                                                                            22;
                                                                                                                                                       Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                   agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                 is not removed.
                                                                                                                                                                                                                INTERLEUKIN-1 BETA; A7CD59EBAC120EC7
                                                                                                                                                                                                                                            n; Inflammatory response;
BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Craniata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
                                                                                                                                            Mismatches
                                                                                                                                                       127;
No. 6.
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i; Cercopithecidae;
                                                                                                                                                       DB 1;
.5e-06;
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                                                                                                                                            61;
                                                                                                                                                                                                                                                                                                                                                                                                                               Usage
                                                                                                                                                                                                                  CRC64
                                                                                                                                                                     Length 269;
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                                                                                   -DKIPVALG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            collaboration -
L outstation -
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RESULT 15 IL1B_MACFA

IL1B_MACFA P79182;

STANDARD;

PRT;

268

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237

YISTSQAENMPVFL-----GGTRGGQDITDFTMQ 265

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Best Local S
Matches 46
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PRINTS; pro0264; INTERLEUKIN1.

PROSITE; pro0263; INTERLEUKIN_1; 1.

PROSITE; pro0263; INTERLEUKIN_1; 1.

Cytokine; Macrophage; Mitogen; Inflammatory response; Pyrogen.

Cytokine; Macrophage; Mitogen; Inflammatory response; Pyrogen.

PROPEP 1116 116 INTERLEUKIN-1 BETA.

20425 MW; CFB7266B3E2C05B4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Totsuka K., Takakura H., Hashimoto O., Tatsumi M.;
Submitted (JUL-1995) to the EMBL/GenBank/DDBJ databases.
Submitted (JUL-1995) to the EMBL/GenBank/DDBJ databases.
FUNCTION: PRODUCED BY ACTIVATED MACROPHAGES, IL-1 STIMULATES
THYMOCYTE PROLIFERATION BY INDUCING IL-2 RELEASE, B-CELL
MATURATION & PROLIFERATION, & FIBROBLAST GROWTH FACTOR ACTIVITY.
IL-1 PROTEINS ARE INVOLVED IN THE INFLAMMATORY RESPONSE, BEING
IDENTIFIED AS ENDOGENOUS PYROGENS, AND ARE REPORTED TO STIMULATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Interpro; IPR000975; ...
Interpro; IPR002348; -..
Pfam; PF00340; interleukin-1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; D63353; BAA09677.1; -. HSSP; P01584; 1HIB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Macaca fascicularis (Crab eating macaque) (Cynomolgus monkey).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE-Thymus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9541;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
120 FLCTVPEADQPVRLTQLPENGGWNA--PITDFYFQ 152 :: | | | | | | | | | | | | | | |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INTERLEUKIN-1 BETA PRECURSOR (IL-1 BETA).
                                                         178
                                                                                                                                        134
                                                                                Local Similarity
nes 46; Conserv
                                                                                                                                                          1 MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILG 60 :|:|| : ::||| : || || ||
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DOMAIN: THE SIMILARITY AMONG THE IL-1 PRECURSORS SUGGESTS THAT THE AMINO ENDS OF THESE PROTEINS SERVE SOME AS YET UNDEFINED FUNCTION.

MISCELLANEOUS: THE LACK OF A SPECIFIC HYDROPHOBIC SEGMENT IN THE PRECURSOR SEQUENCE SUGGESTS THAT IL-1 IS RELEASED BY DAMAGED CELLS OR IS SECRETED BY A MECHANISM DIFFERING FROM THAT USED FOR OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SIMILARITY: BELONGS TO THE IL-1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SECRETORY PROTEINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 THE RELEASE OF PROSTAGLANDIN AND COLLAGENASE FROM SYNOVIAL CELLS
                                                       LKAKNLYLSCYLKDDKPTLQLESVD-PKNYPKKKMEKREVFNKIEINNKLEFESAQFPNW 236
                                                                                                                                    LVMSGPY-----ELKALHLQGQDLEQQVVFSMSFVQGEESN------DKIPVALG 177
                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                     15.3%;
                                                                                                                                                                                                                    21;
                                                                                                                                                                                                                Score 126; DB 1;
Pred. No. 8.2e-06;
11; Mismatches 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   as its content is in
                                                                                                                                                                                                                    62;
                                                                                                                                                                                                                                                       Length 268;
                                                                                                                                                                                                                  Indels
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no
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В 237 YISTSQAESMPVFL-----GTRGGQDITDFTMQ 265

Search completed: Job time: 146 sec June 25, 2001, 14:07:54

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Title:
Perfect score:
Sequence:
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No.
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Listing first 45 summaries
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Maximum DB
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Match Length DB
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Gapop 10.0 , Gapext 0.5
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823
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd
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sp_bacteria:*
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                    Oggyyl mus musculu
Ogjig2 mus musculu
Ogupci homo sapien
Ol4628 homo sapien
Ol4628 homo sapien
Ol4628 tursiops tr
Ogmz4 tursiops tr
Ogmz4 tursiops tr
Oguk2 canis famil
Oguk5 canis famil
Oguk6 homo sapien
Ol7771 eguus cabal
Ogila2 mus musculu
Oguh6 homo sapien
Oguh6 homo sapien
Ognzh6 homo sapien
                                                                                                                                                                            Q9ubhO homo sapien
Q9qyy1 mus musculu
Q9j1g2 mus musculu
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ALIGNMENTS

RN	3 급	R R	RP	2 2	잡	RT	RA	RC	RP S	2 2	RT	RA	RA	RX	ŖΡ	RN	RL	RT	RΑ	RΑ	RX	RP	RN	õ	8	8	20	GN	DE	DE	D	ij	ဌ	AC	IDOO	09119	RESULT
Submitted (JAN-2000) to the EMBL/Genbank/DDBJ databases. [5]	ter locks IL-1, IL-1ra, IL-18 and IL-	10	SEQUENCE FROM N.A.	Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databases.	IL-1ra, IL-18 and IL-18ra activities.	"A tissue specific interleukin-1 receptor antagonist homolog from the	Barton J.L., di Giovine F.S., Symons J.A., Nicklin M.J.H.;		SEQUENCE FROM N.A.	12) 21	A Novel	R., Ford J.E.;	Mulero J.J., Pace A.M., Nelken S.T., Loeb D.B., Correa T.R.,	MEDLINE=99443727; PubMed=10512743;	SEQUENCE FROM N.A.		J. Biol. Chem. 275:1169-1175(2000).	"Four New Members Expand the IL-1 Superfamily.";		Smith D.E., Renshaw B.R., Ketchem R.R., Kubin M., Garka K.E.,		SEQUENCE FROM N.A.	[1]		Primates; Catarrhini; Hominidae;	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	Homo sapiens (Human).		HOMOLOG 1) (INTERLEUKIN-1 DELTA)	(INTERLEUKIN-1 LI	(TrEMBLrel, 16, Last	(TrEMBLrel. 13,	01-MAY-2000 (TrEMBLrel. 13, Created)		O9UBHO PRELIMINARY: PRT: 155 AA.	HO I	

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Q9hbf2 h Q9pvz5

7 homo sapien
5 homo sapien
8 homo sapien
9 homo sapien
9 homo sapien
10 gallus gall
10 homo sapien

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RESULT
OPQYY1
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OPPORTUGE
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Best Local Similarity
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EMBL; AF186094; AAF02757.1; --
EMBL; AJ242737; CAB59822.1; --
EMBL; AJ242738; CAB59823.1; --
EMBL; AJ242738; CAB59823.1; --
EMBL; AJ271336; CAB67704.1; --
EMBL; AF216693; AAF76981.1; --
EMBL; AF230377; AAF91274.1; --
HSSP; P18510; IILR,
HSSP; P18510; IILR
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Q9QYY1;
01-MAY-2000
                                                                                                                                                                                             Barton J.L., Nicklin M.J.H.;
"IL-IL1: A Novel Member of the Interleukin-1 Gene Family
in Trophoblasts and Macrophages.";
Submitted (CCT-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                            MEDLINE=20209405; PubMed=10744718;
Kumar S., McDonnell P.C., Lehr R., Tierney
Griswold D.E., Capper E.A., Tal-Singer R.,
                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
Barton J.L., Nicklin M.J.H.;
                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
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PROSITE; PS00253; INTERLEUKIN_1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Novel IL-1 ligands IL-1d au
Submitted (FEB-2000) to the
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Kastelein R.A.;
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Mulero J.J., Nelken S.T., Ford J.E.;
"Organization of the Human Interleukin-l Receptor Antagonist Gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A. MEDLINE=20322477;
                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                        Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SMART; SM00125; IL1;
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                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID-10090;
                        'Identification and initial characterization the interleukin-1 family.";
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                                                                                                                                                                                                                                                                                                                                                                    musculus (Mouse).
aryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
aryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
malia: Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MVLSGALCFRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSPVILG
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  1-1 family.";
275:10308-10314(2000)
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Pred. No. 3.3e-76;
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                                                                                              L., Tzimas M.
Wells G.I., I
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O9JIG2;
O1-OCT-2000 (TrEMBLrel. 1
01-OCT-2000 (TrEMBLrel. 1
01-MAR-2001 (TrEMBLrel. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Debets R., Timans J.C., Zura
Kastelein R.A.;
"Novel IL-1 ligands IL-1d ar
Submitted (FEB-2000) to the
EMBL; AF230378; AAF91275.1;
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EMBL; AF200495; AAF69251.1;
HSSP; P18510; IILR
InterPro; IPR000975; ...
Pfam; PF00340; IL1; 1.
                                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR000975; -.
pfam; pF00340; ILI; 1.
pRINTS; pR00264; INTERLEUKIN1.
pROSITE; pS00253; INTERLEUKIN_1;
SMART; SM00125; ILI; 1.
SEQUENCE 156 AA; 17136 MW; A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRINTS; PRO0264; INTERLEUKIN1.
PROSITE; PS00253; INTERLEUKIN_1;
SMART; SMO0125; IL1; 1
SEQUENCE 155 AA; 17004 MW; A4
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                          LCTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD
                                                                                       VQGGSQCLSCGVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWF 120
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LCTSPEADQPVRLTQIPEDPAWDAPITDFYFQQCD
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16,
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Last annotation update)
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                                                                                                                                                                                                                                                                               Score 734; DB 11;
Pred. No. 3.8e-67;
4; Mismatches 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 734; DB 11;
Pred. No. 3.8e-67;
4; Mismatches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Craniata; Vertebrata; I
Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IL-le
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    use
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156
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                                                                                                                                                                                                                                                                                                                           Length 156;
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Best Local S
Matches 73
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Q14628;
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01-MAY-2000
01-MAY-2000
01-MAR-2001
SEQUENCE FROM N.A.
MEDLINE=97146044; PubMed=8992991;
Jenkins J.K., Drong R.F., Shuck M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Receptor.
NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                   01-NOV-1996 (TrEMBLrel 01, 01-NOV-1996 (TrEMBLrel 01, 01-MAR-2001 (TrEMBLrel 16, INTRACELLULAR IL-1 RECEPTOR
                                                                                  "Cloning and characterization receptor antagonist.";
J. Exp. Med. 182:623-628(1995)
                                                                                                                                                                                                                                                                Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A. MEDLINE=98183404; PubMed=9514884;
                                                                                                                                             Muzio M., Polentarutti N., Sironi
Introna M., Mantovani A., Colotta
                                                                                                                                                                                      SEQUENCE FROM N.A. MEDLINE=95355865; PubMed=7629520;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pfam; PF00340; IL1; 1.
PRINTS; PR00264; INTERLEUKIN1.
PROSITE; PS00253; INTERLEUKIN_1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SMART; SM00125; IL1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL; AF043143; AAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Weissbach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q9UPC0
                                                                                                                                                                                                                                                 NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                    IL-1RN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Biochem. Biophys. Res. Commun.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 variant."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Weissbach L., Tran K., Colquhoun S.A., Champliaud Detection of an interleukin-1 intracellular recep
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            117
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EADQPVSLTNMPDEG---VMVTKFYFQE 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EADQPVRLTQLPENGGWNAPITDFYFQQ 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLSC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVP 125
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
73; Conserv
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143 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRELIMINARY;
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Pred. No. 5.6e
13; Mismatches
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Last annotation updat
ANTAGONIST TYPE II.
                                                                                                                                                                                                                                                                                                                                                                                                              Created)
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Last annotation update)
RECEPTOR ANTAGONIST VARIANT (FRAGMENT).
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                                                                                                                                                                                                                                                                  Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
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    M.E.,
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    .J.,
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    .L.,
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RESULT
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Query Match
Best Local S
Matches 71
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Best Local
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Pfam; PF00340; ILI; I
PRINTS; PR00254; INTERLEUKIN1.
PRODOm; PD002536; -; 1.
PROSITE; PS00253; INTERLEUKIN_1; 1.
SMART; SM00125; ILI; 1.
SEQUENCE 180 AA; 19897 MW; 624A15
                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=FVBXDBA/1 LACJ;

MEDLINE=98209757; pubMed=9550387;

Gabay C., Porter B., Fantuzzi G., Arend W.

"Mouse IL-1 receptor antagonist isoforms:
and protein expression of intracellular is
distribution of secreted and intracellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INTRACEDLUDAN
Mus musculus (Mouse).
Mus musculus (Mouse).
'Parvota; Metazoa; Chordata; Metazoa; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             070207 PRELIMINARY;
070207;
01-AUG-1998 (TYEMBLIEL 07,
01-AUG-1998 (TYEMBLIEL 07,
01-MAR-2001 (TYEMBLIEL 16,
                                                                                                                          PROSITE; PS00253; INTERLEUKIN_1; SMART; SM00125; IL1; 1. SEQUENCE 159 AA; 17995 MW; A
                                                                                                                                                                                          Pfam; PF00340; IL1; 1.
PRINTS; PR00264; INTERLEUKIN1.
PRODOM; PD002536; -; 1.
                                                                                                                                                                                                                                                                                             J. Immunol. 159:5905-5913(1997).
EMBL; AF001795; AAC15251.1; -.
HSSP; P18510; 1IRA.
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J. Immunol. 158:748-755(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                      InterPro; IPR000975; -.
                                                                                                                                                                                                                                                                                                                                                                         vivo."
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Intracellular IL-1 receptor antagonist
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EADQPVSLTNMPDEG - - - VMVTKFYFQE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLSCVKSGDETRLQLEAVNITDLSENRKQDKRFAFIRSDSGPTTSFESAACPGWFLCTAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FRIWDVNQKTFYLRNNQLVAGYLQGPNVNLEEKIDVVP-----IEPHALFLGIHGGKM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP--VILGVQGGSQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          38.8%;
                         37.5%;
                                                                                                                                                                                                                                                                                                                                                                                           intracellular isoform and and intracellular IL-1 rece
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Last sequence update)
Last annotation update)
RECEPTOR ANTAGONIST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Created)
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                         Score
Pred.
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Pred. No. 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRT;
                                                                                                                          A7CD35D137846222 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             624A1574C2334229 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arend W.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           178
                         308.5;
No. 8.4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     159
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                                                                                                                                                                                                                                                                                                                                                                                                                     complementary DNA cloning soform and tissue
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      180;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     93
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Similarity

Conservative

15;

Mismatches

49;

Indels

13;

Gaps

4.

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DB 11;

Length

159;

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RESULT
OPGMZA
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OPERING
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                                                                                                                                                  Q9GKK2 PRELIMINARY; PRT; 176 AA.
Q9GKK2;
Q9GKK2;
101-MAR-2001 (TrEMBLrel. 16, Created)
101-MAR-2001 (TrEMBLrel. 16, Last sequence update)
101-MAR-2001 (TrEMBLrel. 16, Last annotation update)
1NTERLEUKIN-1 RECEPTOR ANTAGONIST.
Canis familiaris (Dog).
EUKaryota; Metazoa; Chordata; Craniata; Vertebrata; Eu Mammalia; Eutheria; Carnivora; Fissipedia; Canidae; Ca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
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SEQUENCE
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SEQUENCE FROM N.A.

Campbell S.E., Nasir L., Argyle D.J., Gault E., B
"Canine II-1 Receptor Antagonist cDNA Sequence.";
Submitted (DEC-1999) to the EMBL/GenBank/DDBJ dat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tursiops truncatus (Atlantic bottle-nosed dolphin).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Cetacea; Odontoceti; Delphin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Inoue Y., Itou T., Sakai T.;
"Cloning and Sequencing of a Bottle-Nosed Dolphin Interleukin-1 Receptor Antagonist.";
Submitted (FEB-2000) to the EMBL/GenBank/DDBJ databases.
EMBL, AB038268; BAB11806.1;
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SEQUENCE FROM N.A.
Thou T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=9739;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLSC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVP 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ETDQPVGLTNTPQDA---VQVTKFYFQQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP--VILGVQGGSQ 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DRPVSLTNTPEE----PLIVTKFYFQE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FRIWDTNQKTFYLRNNQLIAGYLQGPNIKLEEKIDMVP-----IDLHSVFLGIHGGKLCL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLACVKSGDEIKLGLEPVNITDLNSSKEEDKRFAFIRSDSGPTTSFESAACPGWFLCTAL 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FRIWDVNQKTFYLRNNQLVAGYLQGPNTKLEEKIDVVP-----IEPHAMFLGIHGGKL 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SCAKSGDDIKLQLEEVNITDLSKNKEEDKRFTFIRSEKGPTTSFESAACPGWFLCTTLEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   177 AA;
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48.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Score 305.5; DB 6; Pred. No. 1.9e-23; 13; Mismatches 51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   51;
            databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CRC64;
                                                                      Bennett
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                                                                                                                                                                                       Euteleostomi; Canis.
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                                                                RESULT 10
077771
ID 077771
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Best Local S
Matches 68
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      077771;
077771;
01-NOV-1998
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                                                                                                                                                                                                                   128
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         (TrEMBLrel.
                                                                   PRELIMINARY;
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Created)

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Q9UHA5; PRELIMINARY;
Q9UHA5;
01-MAY-2000 (TrEMBLrel. 1
01-MAY-2000 (TrEMBLrel. 1
01-MAR-2001 (TrEMBLrel. 1
                                                                                                                                                                                                                                                                                                                                                                                                       Sims J.E.;
"Four New Members Expand the IL-1 Superfamily.";
J. Biol. Chem. 275:1169-1175(2000).
EMBL; AP201833; AAF25213.1; -.
HSSP; P10749; 2MIB.
HSSP; P10749; 2MIB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL; AF216526; AAG36777.1; Receptor. SEQUENCE 176 AA; 19938 MV
                                                                                                                                                                                                                                                                                                                Interpro; ipr000975; -.

Pfam; PF00340; IL1; 1.

PRINTS; PR00264; INTERLEUKINI.

PROSITE; PS00253; INTERLEUKIN_1; UNKNOWN_1.

SMART; SM00125; IL1; 1.

SEQUENCE 157 AA; 17702 MW; 7A54F3D7557A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FIL1 ETA.
Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=20092888; PubMed=10625660; Smith D.E., Renshaw B.R., Ketchem
                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
ATSTTSGQPIFLTK -- ERGITNN -- TNFYLDSVE
                                  CTVPEADQPVRLTQLPENGGWNAPITDFYFQQCD 155
                                                                        GKDLCLFCAETQGKPTLQLKEKNIMDLYVEKKAQKPFLFFHNKEGSTSVFQSVSYPGWFI 127
                                                                                         GGSQCLSCG-VGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFL 121
                                                                                                                                                YAIRDSRQMVWVLSGNSLIAAPL--SRSIKPVTLHLIACR--DTEFSDKEKGNMVYLGIK 67
                                                                                                                                                                                  FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP-----VILGVQ 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EADQPVRLTQLPENGGWNAPITDFYFQQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLACVKSGDETRLQLEAVNITDLSKNKDQDKRFTFILSDSGPTTSFESAACPGWFLCTAL 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLSC-GVGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFESAAYPGWFLCTVP 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASLSP--VILGVQGGSQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EADRLVSLTNRPEEA---MMVTKFYFOK 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FRIWDVNQKTFYLRNNQLVAGYLQGSNTKLEEKLDVVP-----VEPHAVFLGIHGGKL 90
                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                              17702 MW; 7A54F3D7557A3EE3 CRC64;
                                                                                                                                                                                                                                        24.4%;
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16,
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Last sequence update)
Last annotation updat
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Pred. No. 5.3e
13; Mismatches
                                                                                                                                                                                                                      Score 200.5; DB 4
Pred. No. 8.2e-13;
4; Mismatches 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8486CA54A254206B
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                                                                                                                                                                                                                                                         DB 4;
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RESULT
Q9JLA2
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Best Local
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01-OCT-2000
01-MAR-2001
        PRINTS;
SMART; S
                                                             Debets R., Timans J., Zurawski S., Bazan J.F., Kastelein R.A "Novel IL-1 family member IL-le responds through the orphan related protein 2; response is antagonized by IL-ld."; Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases. EMBL; AF200493; AAF69249.1; -. EMBL; AF200657; AAG35671.1; -. INTERPROJECTION OF THE PROJECTION OF THE 
                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
MEDLINE=20209405; PubMed=10744718;
Kumar S., McDonnell P.C., Lehr R., Tier
Griswold D.E., Capper E.A., Tal-Singer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Molecular Characterization of Equine Interleukin 1 Rece
Antagonist (IL-Ira) Gene.";
Submitted (JUN-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF072476; AAC62237.1; -.
                                            InterPro; IPR000975; Pfam; PF00340; IL1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=9796;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-OCT-2000 (TremBLrel.
01-OCT-2000 (TremBLrel.
01-MAR-2001 (TremBLrel.
INTERLEUKIN-1 HOMOLOG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q9JLA2;
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PROSITE; PS00253; INTERLEUKIN_1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR000975; -. Pfam; PF00340; IL1; 1.
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Dhar A.K., Thompson M.S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-1998
01-MAR-2001
                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                       "Identification and initial characterization the interleukin-1 family.";
                                                                                                                                                                                                                                                                                                               Young
                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus (Mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Equus caballus (Horse).
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                                                                                                                                                                                                                                                      Biol.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EAVNITDLSKNKEENKRFTFIRSNSGPTTSFESAACPGWFLCTAQEADRPVSLTNKPKE- 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SM00125; IL1;
F0034., IN.... PR00264; IN....
            SM00125;
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(TrEMBLrel.
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                             INTERLEUKIN1
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chordata;
Rodentia;
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Perissodactyla; Equidae;
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15, Last sequence update)
16, Last annotation update)
(INTERLEUKIN-1 EPSILON).
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16, Last annotation update)
ANTAGONIST SECRETORY FORM (F
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Pred. No. 4.2e-1
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                              , Tierney
inger R.,
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Wells G.I., [
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                                                                                                                                                                                                                                                                                        novel members
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RESULT 13
Q9UHA6
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Q9UHA7
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Best Local
Oliman).
FILL ZETA.
HOMO saplens (Human).
Bukaryota; Metazoa; CP
"--mmalia; Eutheria; P
                                                                                                                                                                                                                      Matches
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                                                                  Q9UHA6;
Q9UHA6;
Q1-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q9UHA7;
Q9UHA7;
01-MAY-2000
01-MAY-2000
01-MAR-2001
                                                                                                                                                                                                                                                                      PRINTS; PRO0264; INTERLEUKIN1.
SMART; SM00125; IL1; 1.
SEQUENCE 158 AA; 17684 MW;
                                                                                                                                                                                                                                                                                                                        J. Biol. Chem. 275:1169-1175(2000). EMBL; AF201831; AAF25211.1; -. HSSP; P18510; 1IRA.
                                                01-MAY-2000
01-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
MEDLINE=20092888; PubMed=10625660;
Smith D.E., Renshaw B.R., Ketchem
                                                                                                                                                                                                                                                                                                    InterPro; IPR000975; .
Pfam; PF00340; IL1; 1.
                                                                                                                                                                                                                                                                                                                                                                Sims J.E
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         FIL1 EPSILON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                     "Four New Members Expand the IL-1 Superfamily.";
                                                                                                                                                    114 AAYPGWFLCTVPEADQPVRLTQLPENGGWNAPITDF 149
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12
                                                                                                                                                                                           SPVILGVQGGSQCLSCG-VGQEPTLTLEPVNIMELYLGAKESKSFTFYRRDMGLTSSFES 113
                                                                                                                                        VAFPGWFIAVSSEGGCPLILTQ -- ELG -- KANTTDF 152
                                                                                                                                                                              NPIYLGLNGLNLCLMCAKVGDQPTLQLKEKDIMDLYNQPEPVKSFLFYHSQSGRNSTFES 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IAVCSKGSCPLILTQ -- ELG -- EIFITDF
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                                                                                                                                                                                                                      43;
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                                                (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
                                                                                                                                                                                                                      Conservative
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                                                                                       PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                             (Human)
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Primates;
          Primates;
                  Chordata;
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36.2%;
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                                                Last
Last
                                                                   Created)
                                                                                                                                                                                                                    Score 185.5;
Pred. No. 2.8e
13; Mismatches
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Pred. No. 2.2e
24; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Created)
         Craniata; V
Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                         Craniata; Vo
Catarrhini;
                                                                                       PRT;
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                                                                                                                                                                                                                                                                       469AC84306B0E280 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA0434D68FF62F4A CRC64;
                                                sequence u annotation
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                  Vertebrata; Euteleostomi;
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i; Hominidae;
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                                               on update)
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          Hominidae;
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PRESULT Q9NZH6
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Best Local S
Matches 51
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Sims J.E.;
Four New Members Expand the IL-1 Superfamily.";
J. Biol. Chem. 275:1169-1175(2000).
EMBL; AF001832; AARF25212.1;
HSSP; P18510; IITN.
TTDANNOGTS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    O9NZH6 PRELIMINARY; PRT; 218 AA.
O9NZH6;
O9NZH6;
O1-OCT-2000 (TrEMBLrel. 15, Created)
O1-OCT-2000 (TrEMBLrel. 15, Last sequence update)
O1-MAR-2001 (TrEMBLrel. 16, Last annotation update)
INTERLEUKIN-1 HOMOLOG 4 (IL-1X PROTEIN) (INTERLEUKIN-1-RELATED PROTEIN
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SEQUENCE FROM N.A.
MEDLINE-20092888; PubMed-10625660;
Smith D.E., Renshaw B.R., Ketchem
                                            Submitted (APR-2000) to the EMBL; AF200496; AAF69252.1; EMBL; AF167368; AAG29344.1; EMBL; AF251118; AAG14420.1;
                                                                                                                                                                           Pan G., Risser P., Mao W., Baldwin D.T., Zhong A.W., Yansura Lewis L., Eigenbrot C., Henzle W.J., Vandlen R., Filvaroff E. "IL-1H, an interleukin-1-related protein that binds IL-18 rec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
TISSUE-FETAL LUNG, FETAL TESTIS, FETAL B-CI
TISSUE-FETAL LUNG, FETAL TESTIS, FETAL B-CI
MEDLINE-20209405; PubMed-10744718;
Kumar S., McDonnell P.C., Lehr R., Tierney
Griswold D.E., Capper E.A., Tal-Singer R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR000975; -.
InterPro; IPR000975; -.
Pfam; PF00340; IL1; 1.
PRINTS; PR00264; INTERLEUKIN1
SMART; SM00125; IL1; 1.
                                                                                                                                                                                                                                                                                                                Manoj P.P., Mantovani A., Muzio M.;
Submitted (JUL-1999) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the
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Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                          lRrp."
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J. Biol. Chem. 275:10308-10314(2000).
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PF00340;
                      IPR000975; -.
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36.7%;
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pred. No. 7.4e-10;
"" amatches 56;
                                                                                                                               EMBL/GenBank/DDBJ databases
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Wells G.I., Doy
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Best Local S
Matches 51
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Matches 51
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Q9HBF3; O1-MAR-2001 (TrEMBLrel. 16, Created)
O1-MAR-2001 (TrEMBLrel. 16, Last sequence update)
O1-MAR-2001 (TrEMBLrel. 16, Last annotation updat
INTERLEUKIN-1-RELATED PROTEIN LONG ISOFORM.
                                                                                                                                                                                                                                           Submitted (APR-2000) to the EMBL/GenBank/DDBJ databases EMBL; AF251119; AAG14421.1; . . SEQUENCE 218 AA; 24138 MW; 76E09C35093DEA63 CRC64;
                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.

Pan G., Risser P., Mao W., Baldwin D.T., Zhong A.W., Yansura Lewis L., Eigenbrot C., Henzle W.J., Vandlen R., Filvaroff E.

"IL-1H, an interleukin-1-related protein that binds IL-18 rec
1Rrp.";
                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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SEQUENCE 218 AA; 24126 MW;
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                                                                          QCLSC--GVGQ-EPTLTLEPVNIMELYLGAKES--KSFTFYRRDMGLTSSFESAAYPGWF 120
                                                                                                                FSIHDQDHKVLVLDSGNLIA--VPDKNYIRPEIFFALASSLSSASAEKGSPILLGVSKGE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FRMKDSALKVLYLHNNQLLAGGLHAGKVIKGEEISVVPNRWLDASL---SPVILGVQGGS
ICTSCNCNEPVGVTDKFEN
                            LCTVPEADQPVRLTQLPEN
                                                        FCLYCDKDKGQSHPSLQLKKEKLMKL-AAQKESARRPF1FYRAQVGSWNMLESAAHPGWF
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Similarity 36.7%;
51; Conservative 2:
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Pred. No. 8.7e
21; Mismatches
                                                                                                                                                                                     Score 172.5; DB 4; Length 218; Pred. No. 8.7e-10;
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